SEQUENCE LISTING

<110> Watson, James D.

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KODUKAN'U CAHBOU
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Met Xaa Pro Val Pro Val Ala Thr Ala Ala Tyr
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Val Ser Leu Ala Pro Gly Val Pro Xaa Val Phe Glu Thr
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Gly Gly Gln Ala Ala
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Met Pro Ile Leu Gln Val Ser Gln Thr Gly Arg
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                      5
           <210> 9 .
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           <400> 9
roosis, eyateor
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H
DUSISH
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M
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                                          10
N
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Ala Xaa Val Val Pro Pro Xaa Gly Pro Pro Ala Pro Gly Ala Xaa
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Ala Pro Ala Pro Asp Leu Gln Gly Pro Leu Val Ser Thr Leu Ser
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Ala Thr Pro Asp Trp Ser Gly Arg Tyr Thr Val Val Thr Phe Ala Ser
                5
Asp Lys Leu Gly Thr Ser Val Ala Ala
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Ala Ser Pro Pro Thr Leu Xaa Val Val
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Glu Pro Glu Gly Val Ala Pro Pro
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COSIET ETELEC
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                                    10
Ala Val Asp Pro Xaa Xaa Tyr Val Val
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                                     10
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Phe Ser Arg Pro Gly Leu Pro Val Glu Tyr Leu Met Val Pro Ser
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Phe Ser Arg Pro Gly Leu Pro Val Glu Tyr Leu Met Val Pro Ser Pro
                                     10
Ser Met Gly
      <210> 24
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      <213> Mycobacterium vaccae
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Phe Ser Arg Pro Gly Leu Pro Val Glu Tyr Leu Asp Val Phe Ser
      <210> 25
      <211> 14
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LOUSICH'S CIISOE
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                                     10
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Val Pro Ala Asp Pro Val Gly Ala Ala Gln Ala Glu Pro Ala Xaa
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Xaa Arg Ile Asp
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                 5
      <210> 28
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Ala Pro Ser Leu Ser Val Ser Asp Tyr Ala Arg Asp Ala Gly Phe
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Met Lys Phe Val Asp Arg Phe Arg Gly Ala Val Ala Gly Met Leu Arg
Arg Leu Val Val Glu Ala Met Gly Val Ala Leu Leu Ser Ala Leu Ile
                                25
Gly Val Val Gly Ser Ala Pro Ala Glu Ala Phe Ser Arg Pro Gly Leu
                            40
Pro Val Glu Tyr Leu Gln Val Pro Ser Pro Ser Met Gly Arg Asp Ile
                        55
Lys Val Gln Phe Gln Asn Gly Gly Ala Asn Ser Pro Ala Leu Tyr Leu
Leu Asp Gly Leu Arg Ala Gln Asp Asp Phe Ser Gly Trp Asp Ile Asn
Thr Thr Ala Phe Glu Trp Tyr Tyr Gln Ser Gly Ile Ser Val Val Met
                                105
Pro Val Gly Gln Ser Ser Phe Tyr Ser Asp Trp Tyr Ser Pro Ala
                            120
Cys Gly Lys Ala Gly Cys Gln Thr Tyr Lys Trp Glu Thr Phe Leu Thr
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135
                                            140
    130
Ser Glu Leu Pro Glu Tyr Leu Gln Ser Asn Lys Gln Ile Lys Pro Thr
                    150
                                        155
Gly Ser Ala Ala Val Gly Leu Ser Met Ala Gly Leu Ser Ala Leu Thr
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Leu Ala Ile Tyr His Pro Asp Gln Phe Ile Tyr Val Gly Ser Met Ser
                               185
           180
Gly Leu Leu Asp Pro Ser Asn Ala Met Gly Pro Ser Leu Ile Gly Leu
                           200
Ala Met Gly Asp Ala Gly Gly Tyr Lys Ala Ala Asp Met Trp Gly Pro
                        215
                                            220
Ser Thr Asp Pro Ala Trp Lys Arg Asn Asp Pro Thr Val Asn Val Gly
                    230
                                        235
Thr Leu Ile Ala Asn Asn Thr Arg Ile Trp Met Tyr Cys Gly Asn Gly
                                    250
Lys Pro Thr Glu Leu Gly Gly Asn Asn Leu Pro Ala Lys Leu Leu Glu
            260
                               265
Gly Leu Val Arg Thr Ser Asn Ile Lys Phe Gln Asp Gly Tyr Asn Ala
                            280
Gly Gly Gly His Asn Ala Val Phe Asn Phe Pro Asp Ser Gly Thr His
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Ser Trp Glu Tyr Trp Gly Glu Gln Leu Asn Asp Met Lys Pro Asp Leu
                   310
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Gln Gln Tyr Leu Gly Ala Thr Pro Gly Ala
                325
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Val Gly Ala Ala Ala Thr Leu Pro Ser Leu Ile Ser Leu Ala Gly Gly
            20
                                25
Ala Ala Thr Ala Ser Ala Phe Ser Arg Pro Gly Leu Pro Val Glu Tyr
                            40
Leu Gln Val Pro Ser Glu Ala Met Gly Arg Thr Ile Lys Val Gln Phe
Gln Asn Gly Gly Asn Gly Ser Pro Ala Val Tyr Leu Leu Asp Gly Leu
                                        75
                    70
Arg Ala Gln Asp Asp Tyr Asn Gly Trp Asp Ile Asn Thr Ser Ala Phe
Glu Trp Tyr Tyr Gln Ser Gly Leu Ser Val Val Met Pro Val Gly Gly
            100
                               105
Gln Ser Ser Phe Tyr Ser Asp Trp Tyr Ser Pro Ala Cys Gly Lys Ala
                            120
Gly Cys Thr Thr Tyr Lys Trp Glu Thr Phe Leu Thr Ser Glu Leu Pro
                        135
Lys Trp Leu Ser Ala Asn Arg Ser Val Lys Ser Thr Gly Ser Ala Val
                                        155
                    150
Val Gly Leu Ser Met Ala Gly Ser Ser Ala Leu Ile Leu Ala Ala Tyr
                                    170
His Pro Asp Gln Phe Ile Tyr Ala Gly Ser Leu Ser Ala Leu Met Asp
                                185
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Ser Ser Gln Gly Ile Glu Pro Gln Leu Ile Gly Leu Ala Met Gly Asp

195 200 205 Ala Gly Gly Tyr Lys Ala Ala Asp Met Trp Gly Pro Pro Asn Asp Pro 215 Ala Trp Gln Arg Asn Asp Pro Ile Leu Gln Ala Gly Lys Leu Val Ala 230 235 Asn Asn Thr His Leu Trp Val Tyr Cys Gly Asn Gly Thr Pro Ser Glu 250 245 Leu Gly Gly Thr Asn Val Pro Ala Glu Phe Leu Glu Asn Phe Val His 260 265 Gly Ser Asn Leu Lys Phe Gln Asp Ala Tyr Asn Gly Ala Gly Gly His 275 280 Asn Ala Val Phe Asn Leu Asn Ala Asp Gly Thr His Ser Trp Glu Tyr 295 300 Trp Gly Ala Gln Leu Asn Ala Met Lys Pro Asp Leu Gln Asn Thr Leu 310 315 Met Ala Val Pro Arg Ser Gly 325

Met Gln Leu Val Asp Arg Val Arg Gly Ala Val Thr Gly Met Ser Arg

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<212> PRT

<213> Mycobacterium tuberculosis

<400> 32

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	260			265					270		
Glu Gly Phe	Val Arg	Thr Ser	Asn	Ile	Lys	Phe	Gln	Asp	Ala	Tyr	Asn
275	*		280					285			
Ala Gly Gly	Gly His	Asn Gly	val	Phe	Asp	Phe	Pro	Asp	Ser	Gly	Thr
290		295	i				300				
His Ser Trp	Glu Tyr	Trp Gly	Ala	Gln	Leu	Asn	Ala	Met	Lys	Pro	Asp
305		310				315					320
Leu Gln Arg	Ala Leu	Gly Ala	Thr	Pro	Asn	Thr	Gly	Pro	Ala	Pro	Gln
	325				330					335	
Gly Ala											

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<213> Mycobacterium tuberculosis

<400> 33

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Glu Tyr Trp Gly Ala Gln Leu Asn Ala Met Lys Gly Asp Leu Gln Ser

Ser Leu Gly Ala Gly 325 <210> 34 <211> 338 <212> PRT <213> Mycobacterium bovis <400> 34 Met Gln Leu Val Asp Arg Val Arg Gly Ala Val Thr Gly Met Ser Arg Arg Leu Val Val Gly Ala Val Gly Ala Ala Leu Val Ser Gly Leu Val Gly Ala Val Gly Gly Thr Ala Thr Ala Gly Ala Phe Ser Arg Pro Gly 40 Leu Pro Val Glu Tyr Leu Gln Val Pro Ser Pro Ser Met Gly Arg Asp 55 Ile Lys Val Gln Phe Gln Ser Gly Gly Ala Asn Ser Pro Ala Leu Tyr 70 Leu Leu Asp Gly Leu Arg Ala Gln Asp Asp Phe Ser Gly Trp Asp Ile 85 90 Asn Thr Pro Ala Phe Glu Trp Tyr Asp Gln Ser Gly Leu Ser Val Val Met Pro Val Gly Gly Gln Ser Ser Phe Tyr Ser Asp Trp Tyr Gln Pro 120 Ala Cys Gly Lys Ala Gly Cys Gln Thr Tyr Lys Trp Glu Thr Phe Leu 135 Thr Ser Glu Leu Pro Gly Trp Leu Gln Ala Asn Arg His Val Lys Pro 150 155 Thr Gly Ser Ala Val Val Gly Leu Ser Met Ala Ala Ser Ser Ala Leu 170 165 Thr Leu Ala Ile Tyr His Pro Gln Gln Phe Val Tyr Ala Gly Ala Met 180 185 Ser Gly Leu Leu Asp Pro Ser Gln Ala Met Gly Pro Thr Leu Ile Gly 200 205 Leu Ala Met Gly Asp Ala Gly Gly Tyr Lys Ala Ser Asp Met Trp Gly 215 220 Pro Lys Glu Asp Pro Ala Trp Gln Arg Asn Asp Pro Leu Leu Asn Val 235 Gly Lys Leu Ile Ala Asn Asn Thr Arg Val Trp Val Tyr Cys Gly Asn 245 250 Gly Lys Pro Ser Asp Leu Gly Gly Asn Asn Leu Pro Ala Lys Phe Leu 265 Glu Gly Phe Val Arg Thr Ser Asn Ile Lys Phe Gln Asp Ala Tyr Asn 280 Ala Gly Gly Gly His Asn Gly Val Phe Asp Phe Pro Asp Ser Gly Thr 295 300 His Ser Trp Glu Tyr Trp Gly Ala Gln Leu Asn Ala Met Lys Pro Asp Leu Gln Arg Ala Leu Gly Ala Thr Pro Asn Thr Gly Pro Ala Pro Gln 330 Gly Ala

315

320

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Gly Ala Gly

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<211> 333

310

<213> Mycobacterium leprae

<400> 36

<212> PRT

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Asn Ala Val Phe Asn Phe Pro Pro Asn Gly Thr His Ser Trp Glu Tyr

Trp Gly Ala Gln Leu Asn Ala Met Lys Gly Asp Leu Gln Ser Ser Leu

295

Gly Leu Val Gly Val Val Gly Asp Thr Ala Ile Ala Val Ala Phe Ser Lys Pro Gly Leu Pro Val Glu Tyr Leu Gln Val Pro Ser Pro Ser Met 55 Gly His Asp Ile Lys Ile Gln Phe Gln Gly Gly Gln His Ala Val Tyr Leu Leu Asp Gly Leu Arg Ala Gln Glu Asp Tyr Asn Gly Trp Asp Ile Asn Thr Pro Ala Phe Glu Glu Tyr Tyr His Ser Gly Leu Ser Val 100 105 Ile Met Pro Val Gly Gly Gln Ser Ser Phe Tyr Ser Asn Trp Tyr Gln 120 Pro Ser Gln Gly Asn Gly Gln His Tyr Thr Tyr Lys Trp Glu Thr Phe 135 140 Leu Thr Gln Glu Met Pro Ser Trp Leu Gln Ala Asn Lys Asn Val Leu 150 155 Pro Thr Gly Asn Ala Ala Val Gly Leu Ser Met Ser Gly Ser Ser Ala 170 Leu Ile Leu Ala Ser Tyr Tyr Pro Gln Gln Phe Pro Tyr Ala Ala Ser 180 185 Leu Ser Gly Phe Leu Asn Pro Ser Glu Gly Trp Trp Pro Thr Met Ile 195 200 205 Gly Leu Ala Met Asn Asp Ser Gly Gly Tyr Asn Ala Asn Ser Met Trp 215 220 Gly Pro Ser Thr Asp Pro Ala Trp Lys Arg Asn Asp Pro Met Val Gln 230 235 Ile Pro Arg Leu Val Ala Asn Asn Thr Arg Ile Trp Val Tyr Cys Gly 245 250 Asn Gly Ala Pro Asn Glu Leu Gly Gly Asp Asn Ile Pro Ala Lys Phe 265 Leu Glu Ser Leu Thr Leu Ser Thr Asn Glu Ile Phe Gln Asn Thr Tyr 280 Ala Ala Ser Gly Gly Arg Asn Gly Val Phe Asn Phe Pro Pro Asn Gly 295 300 Thr His Ser Trp Pro Tyr Trp Asn Gln Gln Leu Val Ala Met Lys Pro 310 315 Asp Ile Gln Gln Ile Leu Asn Gly Ser Asn Asn Asn Ala 325

<210> 37

<211> 340

<212> PRT

<213> Mycobacterium tuberculosis

<400> 37

 Met
 Thr
 Phe
 Phe
 Glu
 Gln
 Val
 Arg
 Arg
 Leu
 Arg
 Ser
 Ala
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 Thr
 Thr
 Thr
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 Leu
 Pro
 Arg
 Arg
 Val
 Ala
 Ile
 Ala
 Met
 Gly
 Ala
 Val
 Leu
 Val
 Tyr

 Gly
 Leu
 Val
 Gly
 Pro
 Ala
 Thr
 Ala
 Gly
 Ala
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 Arg
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	Ile	Asn	Thr	Pro 100	Ala	Phe	Glu	Glu	Tyr 105	Tyr	Gln	Ser	Gly	Leu 110	Ser	Val	
	Ile	Met	Pro 115	Val	Gly	Gly	Gln	Ser 120	Ser	Phe	Tyr	Thr	Asp 125	Trp	Tyr	Gln	
	Pro	Ser 130	Gln	Ser	Asn	Gly	Gln 135	Asn	Tyr	Thr	Tyr	Lys 140	Trp	Glu	Thr	Phe	
	Leu 145	Thr	Arg	Glu	Met	Pro 150	Ala	Trp	Leu	Gln	Ala 155	Asn	Lys	Gly	Val	Ser 160	
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	Ile	Pro	Arg	Leu	Val 245	Ala	Asn	Asn	Thr	Arg 250	Ile	Trp	Val	Tyr	Cys 255	Gly	
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	Leu	Glu	Gly 275	Leu	Thr	Leu	Arg	Thr 280	Asn	Gln	Thr	Phe	Arg 285	Asp	Thr	Tyr	
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,	305			_		310	_				315				Lys	320	
	_				Val 325	Leu	Asn	Gly	Ala	Thr 330	Pro	Pro	Ala	Ala	Pro 335	Ala	
	Ala	Pro	Ala	Ala 340													
			210> 211>				,										
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	cada		100>	39 cgtto	attaa	ac.											20
	cayo		999 (210>		, c c y <u>c</u>	5 ~											20
				1211	L												

<212> DNA <213> Mycobacterium vaccae

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<212> DNA <213> Mycobacterium vaccae

<400> 41

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<210> 42 <211> 1052 <212> DNA <213> Mycobacterium vaccae

<400> 42

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agcgcggcgc	tgacctacgc	gatccatcac	ccgcagcagt	tcatctacgc	ctcgtcgctg	600
tcaggcttcc	tgaacccgtc	cgagggctgg	tggccgatgc	tgatcgggct	ggcgatgaac	660
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cgcaacgacc	cgatggtcaa	catcaaccag	ctggtggcca	acaacacccg	gatctggatc	780
tactgcggca	ccggcacccc	gt.cggagctg	gacaccggga	ccccgggcca	gaacctgatg	840
gccgcgcagt	tcctcgaagg	attcacgttg	cggaccaaca	tcgccttccg	tgacaactac	900
atcgcagccg	gcggcaccaa	cggtgtcttc	aacttcccgg	cctcgggcac	ccacagctgg	960
gggtactggg	ggcagcagct	gcagcagatg	aagcccgaca	tccagcgggt	tctgggagct	1020
caggccaccg	cctagccacc	caccccacac	CC			1052

<210> 43

<211> 326

<212> PRT

<213> Mycobacterium vaccae

<400> 43

Leu Gln Ala His Leu Leu

Gly Gly Ser Ala Thr Ala Gly Ala Phe Ser Arg Pro Gly Leu Pro Val 40 Glu Tyr Leu Met Val Pro Ser Pro Ser Met Gly Arg Asp Ile Lys Ile Gln Phe Gln Ser Gly Gly Glu Asn Ser Pro Ala Leu Tyr Leu Leu Asp 70 Gly Leu Arg Ala Gln Glu Asp Phe Asn Gly Trp Asp Ile Asn Thr Gln Ala Phe Glu Trp Phe Leu Asp Ser Gly Ile Ser Val Val Met Pro Val 105 Gly Gly Gln Ser Ser Phe Tyr Thr Asp Trp Tyr Ala Pro Ala Arg Asn 120 125 Lys Gly Pro Thr Val Thr Tyr Lys Trp Glu Thr Phe Leu Thr Gln Glu 135 140 Leu Pro Gly Trp Leu Gln Ala Asn Arg Ala Val Lys Pro Thr Gly Ser 150 155 Gly Pro Val Gly Leu Ser Met Ala Gly Ser Ala Ala Leu Asn Leu Ala 170 165 Thr Trp His Pro Glu Gln Phe Ile Tyr Ala Gly Ser Met Ser Gly Phe 185 Leu Asn Pro Ser Glu Gly Trp Trp Pro Phe Leu Ile Asn Ile Ser Met 200 Gly Asp Ala Gly Gly Phe Lys Ala Asp Asp Met Trp Gly Lys Thr Glu 215 220 Gly Ile Pro Thr Ala Val Gly Gln Arg Asn Asp Pro Met Leu Asn Ile 230 235 Pro Thr Leu Val Ala Asn Asn Thr Arg Ile Trp Val Tyr Cys Gly Asn 245 250 Gly Gln Pro Thr Glu Leu Gly Gly Gly Asp Leu Pro Ala Thr Phe Leu 265 Glu Gly Leu Thr Ile Arg Thr Asn Glu Thr Phe Arg Asp Asn Tyr Ile 280 Ala Ala Gly Gly His Asn Gly Val Phe Asn Phe Pro Ala Asn Gly Thr 295 His Asn Trp Ala Tyr Trp Gly Arg Glu Leu Gln Ala Met Lys Pro Asp 315

Met Arg Leu Leu Asp Arg Ile Arg Gly Pro Trp Ala Arg Arg Phe Gly

Val Val Ala Val Ala Thr Ala Met Met Pro Ala Leu Val Gly Leu Ala 20 25 30

<400> 44

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<210> 44
<211> 161
<212> PRT
<213> Mycobacterium vaccae
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Ser Gly Trp Asp Ile Asn Thr Ala Ala Phe Glu Trp Tyr Val Asp Ser 10 Gly Leu Ala Val Ile Met Pro Val Gly Gly Gln Ser Ser Phe Tyr Ser 20 25 Asp Trp Tyr Ser Pro Ala Cys Gly Lys Ala Gly Cys Gln Thr Tyr Lys 40 Trp Glu Thr Phe Leu Thr Gln Glu Leu Pro Ala Tyr Leu Ala Ala Asn 55 Lys Gly Val Asp Pro Asn Arg Asn Ala Ala Val Gly Leu Ser Met Ala 70 75 Gly Ser Ala Ala Leu Thr Leu Ala Ile Tyr His Pro Gln Gln Phe Gln 90 Tyr Ala Gly Ser Leu Ser Gly Tyr Leu Asn Pro Ser Glu Gly Trp Trp 100 105 Pro Met Leu Ile Asn Ile Ser Met Gly Asp Ala Gly Gly Tyr Lys Ala Asn Asp Met Trp Gly Pro Pro Lys Asp Pro Ser Ser Ala Trp Lys Arg 135 Asn Asp Pro Met Val Asn Ile Gly Lys Leu Val Ala Asn Asn Thr Pro 155 Leu

<210> 45 <211> 334 <212> PRT <213> Mycobacterium vaccae

<400> 45

Met Lys Phe Thr Glu Lys Trp Arg Gly Ser Ala Lys Ala Ala Met His Arg Val Gly Val Ala Asp Met Ala Ala Val Ala Leu Pro Gly Leu Ile Gly Phe Ala Gly Gly Ser Ala Thr Ala Gly Ala Phe Ser Arg Pro Gly 40 Leu Pro Val Glu Tyr Leu Asp Val Phe Ser Pro Ser Met Gly Arg Asp 60 Ile Arg Val Gln Phe Gln Gly Gly Gly Thr His Ala Val Tyr Leu Leu Asp Gly Leu Arg Ala Gln Asp Asp Tyr Asn Gly Trp Asp Ile Asn Thr 85 90 Pro Ala Phe Glu Trp Phe Tyr Glu Ser Gly Leu Ser Thr Ile Met Pro 105 Val Gly Gly Gln Ser Ser Phe Tyr Ser Asp Trp Tyr Gln Pro Ser Arg 120 Gly Asn Gly Gln Asn Tyr Thr Tyr Lys Trp Glu Thr Phe Leu Thr Gln 135 140 Glu Leu Pro Thr Trp Leu Glu Ala Asn Arg Gly Val Ser Arg Thr Gly 155

Asn Ala Phe Val Gly Leu Ser Met Ala Gly Ser Ala Ala Leu Thr Tyr 165 170 Ala Ile His His Pro Gln Gln Phe Ile Tyr Ala Ser Ser Leu Ser Gly 185 Phe Leu Asn Pro Ser Glu Gly Trp Trp Pro Met Leu Ile Gly Leu Ala 200 Met Asn Asp Ala Gly Gly Phe Asn Ala Glu Ser Met Trp Gly Pro Ser 215 Ser Asp Pro Ala Trp Lys Arg Asn Asp Pro Met Val Asn Ile Asn Gln 235 230 Leu Val Ala Asn Asn Thr Arg Ile Trp Ile Tyr Cys Gly Thr Gly Thr 250 245 Pro Ser Glu Leu Asp Thr Gly Thr Pro Gly Gln Asn Leu Met Ala Ala Gln Phe Leu Glu Gly Phe Thr Leu Arg Thr Asn Ile Ala Phe Arg Asp 275 280 Asn Tyr Ile Ala Ala Gly Gly Thr Asn Gly Val Phe Asn Phe Pro Ala 295 300 Ser Gly Thr His Ser Trp Gly Tyr Trp Gly Gln Gln Leu Gln Met 310 315 Lys Pro Asp Ile Gln Arg Val Leu Gly Ala Gln Ala Thr Ala 325 330 <210> 46 <211> 795 <212> DNA <213> Mycobacterium vaccae

<400> 46

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<210> 47 <211> 142 <212> PRT <213> Mycobacterium vaccae

<400> 47

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50
                        55
                                            60
Asn Ala Asp Ala Tyr Ala Phe Ala Lys Arg Glu Lys Val Ser Leu Ala
                    70
                                        75
Pro Gly Val Pro Trp Val Phe Glu Thr Thr Met Ala Asp Pro Asn Trp
Ala Ile Leu Gln Val Ser Ser Thr Thr Arg Gly Gln Ala Ala Pro
            100
                                105
Asn Ala His Cys Asp Ile Ala Val Asp Gly Gln Glu Val Leu Ser Gln
                            120
His Asp Asp Pro Tyr Asn Val Arg Cys Gln Leu Gly Gln Trp
    130
      <210> 48
      <211> 300
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 48
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cgaaggetge gegaegagtt egaetteegg tatetegteg gegtegtete ggtgatgggg
                                                                       120
gacaaggacg tggacgggat ccgccaggac ccgggcgtgc cggacgggcg cggtctcgca
                                                                       180
ctgttcgtct cgggcgacaa ccttcgaaag ggtgcggcgc tcaacacgat ccagatcgcc
                                                                       240
gagetgetgg cegeceagtt gtaagtgtte egeegaaatt geatteeaeg cegataateg
                                                                       300
      <210> 49
      <211> 563
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 49
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                                                                        60
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                                                                       120
gtegagettg ccaaagtgca actgggtgaa ggtttetege acaccaegge egtgttggee
                                                                       180
accepted atgtgctgct caagettetg cacceggtca tgccgttcgt caccgaggtg
                                                                       240
ctgtggaagg ccctgaccgg gcgggccggc gcgagcgaac gtctgggaaa tgtggagtca
                                                                       300
ctggtcgtcg cggactggcc cacgcccacc ggatacgcgc tggatcaggc tgccgcacaa
                                                                       360
cggatcgccg acacccagaa gttgatcacc gaggtgcgcc ggttccgcag cgatcagggt
                                                                       420
ctggccgacc gccagcgggt gcctgcccgg ttgtccggca tcgacaccgc gggtctggac
                                                                       480
gcccatgtcc cggcggtgcg cgcgctggcc tggcttgacc gagggtgatg agggcttcac
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cgcgtccgaa tcggtcgagg tgc
                                                                       563
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      <211> 434
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 50
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geogacgege ceategacgt egeogtegte gaggteggge teggteggteg etgggacgeg
                                                                       120
acgaacgtgg tgaacgcacc ggtcgcggtc atcaccccga tcggggtgga ccacaccgac
                                                                       180
tacctcggtg acacgatcgc cgagatcgcc ggggagaagg ccggaaatca tcacccgcca
                                                                       240
geogacgace tggtgccgac cgacaccgtc geogtgctgg cgcggcaggt teccgaggcc
                                                                       300
atggaggtgc tgctggccca ggcggtgcgc tcggatgcgg ctgtagcgcg cgaggattcg
                                                                       360
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                                                                       420
                                                                       434
tcggtggcgt ctac
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<210> 51

<211> 438 <212> DNA <213> Mycobacterium vaccae ggateceaet eeegegeegg eggeggeeag etggtaegge eatteeageg tgetgatega 60 ggtcgacggc taccgcgtgc tggccgaccc ggtgtggagc aacagatgtt cgccctcacg 120 ggcggtcgga ccgcagcgca tgcacgacgt cccggtgccg ctggaggcgc ttcccgccgt 180 ggacgeggtg gtgategeca acgaccacta egaccacete gacategaca ecategtege 240 gttggcgcac acccageggg ccccgttcgt ggtgccgttg ggcatcggcg cacacctgcg 300 caagtggggc gtccccgagg cgcggatcgt cgagttggac tggcacgaag cccaccgcat 360 egacgacetg aegetggtet geacceeege eeggeactte teeggeeggt tgtteteeeg 420 cgactcgacg ctgtgggc 438 <210> 52 <211> 87 <212> PRT <213> Mycobacterium vaccae <400> 52 Ala Ser Ala Pro Thr Val Phe Ile Asp Ala Ala His Asn Pro Gly Gly 10 5 Pro Cys Ala Cys Arg Arg Leu Arg Asp Glu Phe Asp Phe Arg Tyr Leu 25 Val Gly Val Val Ser Val Met Gly Asp Lys Asp Val Asp Gly Ile Arg 40 Gln Asp Pro Gly Val Pro Asp Gly Arg Gly Leu Ala Leu Phe Val Ser 55 Gly Asp Asn Leu Arg Lys Gly Ala Ala Leu Asn Thr Ile Gln Ile Ala 70 Glu Leu Leu Ala Ala Gln Leu <210> 53 <211> 175 <212> PRT <213> Mycobacterium vaccae <400> 53 Gly Ser Ser Ala Gly Ser Arg Val Arg Ala Glu Val Asp Val Thr Leu 1 5 10 Asp Gly Tyr Glu Phe Ser Arg Ala Cys Glu Ala Leu Tyr His Phe Ala 25 Trp Asp Glu Phe Cys Asp Trp Tyr Val Glu Leu Ala Lys Val Gln Leu 40 Gly Glu Gly Phe Ser His Thr Thr Ala Val Leu Ala Thr Val Leu Asp 55 Val Leu Leu Lys Leu Leu His Pro Val Met Pro Phe Val Thr Glu Val 70 75 Leu Trp Lys Ala Leu Thr Gly Arg Ala Gly Ala Ser Glu Arg Leu Gly 90 Asn Val Glu Ser Leu Val Val Ala Asp Trp Pro Thr Pro Thr Gly Tyr 105 Ala Leu Asp Gln Ala Ala Gln Arg Ile Ala Asp Thr Gln Lys Leu 120 125

140

Ile Thr Glu Val Arg Arg Phe Arg Ser Asp Gln Gly Leu Ala Asp Arg

Trp 145

Gln Arg Val Pro Ala Arg Leu Ser Gly Ile Asp Thr Ala Gly Leu Asp 145 150 155 Ala His Val Pro Ala Val Arg Ala Leu Ala Trp Leu Asp Arg Gly 170 <210> 54 <211> 144 <212> PRT <213> Mycobacterium vaccae <400> 54 Gly Pro Gly Pro Arg Asn Ser Lys Phe Glu Val Val Thr Gly Met Ala Phe Ala Ala Phe Ala Asp Ala Pro Ile Asp Val Ala Val Val Glu Val Gly Leu Gly Gly Arg Trp Asp Ala Thr Asn Val Val Asn Ala Pro Val 40 Ala Val Ile Thr Pro Ile Gly Val Asp His Thr Asp Tyr Leu Gly Asp Thr Ile Ala Glu Ile Ala Gly Glu Lys Ala Gly Asn His His Pro Pro 70 75 Ala Asp Asp Leu Val Pro Thr Asp Thr Val Ala Val Leu Ala Arg Gln 90 Val Pro Glu Ala Asn Glu Val Leu Leu Ala Gln Ala Val Arg Ser Asp 100 105 Ala Ala Val Ala Arg Glu Asp Ser Glu Cys Ala Val Leu Gly Arg Gln 120 Val Ala Ile Gly Gly Ser Cys Ser Gly Cys Arg Gly Ser Val Ala Ser 135 <210> 55 <211> 145 <212> PRT <213> Mycobacterium vaccae <400> 55 Asp Pro Thr Pro Ala Pro Ala Ala Ala Ser Trp Tyr Gly His Ser Ser Val Leu Ile Glu Val Asp Gly Tyr Arg Val Leu Ala Asp Pro Val Trp 25 Ser Asn Arg Cys Ser Pro Ser Arg Ala Val Gly Pro Gln Arg Met His 40 Asp Val Pro Val Pro Leu Glu Ala Leu Pro Ala Val Asp Ala Val Val 55 Ile Ser Asn Asp His Tyr Asp His Leu Asp Ile Asp Thr Ile Val Ala Leu Ala His Thr Gln Arg Ala Pro Phe Val Val Pro Leu Gly Ile Gly 85 90 Ala His Leu Arg Lys Trp Gly Val Pro Glu Ala Arg Ile Val Glu Leu Asp Trp His Glu Ala His Arg Ile Asp Asp Leu Thr Leu Val Cys Thr Pro Ala Arg His Phe Ser Gly Arg Leu Phe Ser Arg Asp Ser Thr Leu

135

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           <211> 10
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           <213> Mycobacterium vaccae
           <220>
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           <223> Residue can be either Gly, Ile, Leu or Val
           <221> UNSURE
           <222> (2)...(2)
           <223> Residue can be either Ile, Leu, Gly, or Ala
           <221> UNSURE
           <222> (5)...(5)
           <221> UNSURE
           <222> (9)...(9)
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    Xaa Xaa Ala Pro Xaa Gly Asp Ala Xaa Arg
-
                      5
           <210> 57
           <211> 8
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          <220>
           <221> UNSURE
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           <223> Residue can be either Ile or Leu
          <400> 57
    Pro Glu Ala Glu Ala Asn Xaa Arg
                      5
           <210> 58
           <211> 11
           <212> PRT
           <213> Mycobacterium vaccae
           <220>
           <221> UNSURE
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           <223> Residue can be either Gln or Gly
          <221> UNSURE
           <222> (5)...(5)
           <223> Residue can be either Gly or Gln
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     1
                      5
           <210> 59
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<211> 34
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                                 25
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      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <400> 60
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                                                                         20
      <210> 61
      <211> 20
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <400> 61
                                                                         20
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      <210> 62
      <211> 313
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 62
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                                                                         60
                                                                        120
qcaqqqaatq tcqcaqqacc cqqtcqcqqt qqcqqcctcq aacaatccgq agttgacaac
gctgtacggc tgcactgtcg ggccagctca atccgcaagt aaacctggtg gacaccctca
                                                                        180
acageggtea gtacaeggtg ttegcaeega ceaaegegge atttageaag etgeeggeat
                                                                        240
ccacqatcga cgagctcaag accaattcgt cactgctgac cagcatcctg acctaccacg
                                                                        300
tggtggccgg cca
                                                                        313
      <210> 63
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      <221> UNSURE
      <222> (7)...(17)
      <400> 63
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Glu Pro Ala Gly Pro Leu Pro Xaa Tyr Asn Glu Arg Leu His Thr Leu
     1
    Xaa Gln
          <210> 64
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          <222> (21)...(21)
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                    5
    Thr Val Gln Gln Xaa Asp Thr Phe Leu
                20
          <210> 65
          <211> 26
          <212> PRT
          <213> Mycobacterium vaccae
          <220>
          <221> UNSURE
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          <222> (21)...(22)
          <221> UNSURE
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          <400> 65
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ΠJ
    Glu Pro Gly Leu Xaa Xaa Val Xaa Asp Ala
                20
          <210> 66
          <211> 32
          <212> DNA
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          <400> 66
                                                                             32
    accgccctcg agttctcccg gccaggtctg cc
          <210> 67
          <211> 32
          <212> DNA
          <213> Artificial Sequence
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<220>
      <223> Made in a lab
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      <210> 68
      <211> 30
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      <220>
      <223> Made in a lab
      <400> 68
catggatcca ttctcccggc ccggtcttcc
                                                                        30
      <210> 69
      <211> 26
      <212> DNA
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      <223> Made in a lab
      <400> 69
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                                                                        26
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      <212> PRT
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Ser Gly Trp Asp Ile Asn Thr Ala Ala Phe Glu Trp Tyr Val Asp Ser
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Gly Leu Ala Val Ile Met Pro Val Gly Gly Gln Ser Ser Phe Tyr Ser
            20
                                25
Asp Trp Tyr Ser Pro Ala Cys Gly Lys Ala Gly Cys Gln Thr Tyr Lys
Trp Glu Thr Phe Leu Thr Gln Glu Leu Pro Ala Tyr Leu Ala Ala Asn
                        55
Lys Gly Val Asp Pro Asn Arg Asn Ala Ala Val Gly Leu Ser Met Ala
                    70
                                        75
Gly Ser Ala Ala Leu Thr Leu Ala Ile Tyr His Pro Gln Gln Phe Gln
                                    90
Tyr Ala Gly Ser Leu Ser Gly Tyr Leu Asn Pro Ser Glu Gly Trp Trp
                                105 .
            100
Pro Met Leu Ile Asn Ile Ser Met Gly Asp Ala Gly Gly Tyr Lys Ala
Asn Asp Met Trp Gly Arg Thr Glu Asp Pro Ser Ser Ala Trp Lys Arg
                        135
                                            140
Asn Asp Pro Met Val Asn Ile Gly Lys Leu Val Ala Asn Asn Thr Pro
                    150
                                        155
Leu
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<220> <223> Made in a lab	
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<210> 72	
<211> 32	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Made in a lab	
<400> 72	
gagagacteg agtgacteae caetgacega ge	32
<210> 73	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Made in a lab	
<221> unsure	
<222> (3)(3)	
<221> unsure	
<222> (6)(6)	
<221> unsure	
<222> (9)(9)	
<221> unsure	
<222> (15)(15)	
<400> 73	
ggngengene argengaree	20
<210> 74	
<211> 825	
<212> DNA	
<213> Mycobacterium vaccae	
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ctgctgaccg ctgccgacgc	cgagcgggta	cgcctgaccg	tgccgattcc	cggtcagcgg	780
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85 90 95 Gly Ala His Leu Arg Lys Trp Gly Val Pro Glu Ala Arg Ile Val Glu

100 105 110 Leu Asp Trp His Glu Ala His Arg Ile Asp Asp Leu Thr Leu Val Cys

\$115\$ \$120\$ \$125\$ Thr Pro Ala Arg His Phe Ser Gly Arg Leu Phe Ser Arg Asp Ser Thr

130 135 140

Leu Trp Ala Ser Trp Val Val Thr Gly Ser Ser His Lys Ala Phe Phe
145 150 150 155 160

Gly Gly Asp Thr Gly Tyr Thr Lys Ser Phe Ala Glu Ile Gly Asp Glu 165 170 175

Tyr Gly Pro Phe Asp Leu Thr Leu Leu Pro Ile Gly Ala Tyr His Pro 180 185 190

Ala Phe Ala Asp Ile His Met Asn Pro Glu Glu Ala Val Arg Ala His 195 200 205

Leu Asp Leu Thr Glu Val Asp Asn Ser Leu Met Val Pro Ile His Trp 210 215 220

Ala Thr Phe Arg Leu Ala Pro His Pro Trp Ser Glu Pro Ala Glu Arg 225 230 235 240

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Phe

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aaggeteegg getteggtga eegeegeaag gegatgetge aggacatgge cateeteace
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Asn Lys Ile Arq Gly Thr Phe Lys Ser Val Ala Val Lys Ala Pro Gly
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Phe Gly Asp Arg Arg Lys Ala Met Leu Gln Asp Met Ala Ile Leu Thr
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gaggetgtea eccagteget getgaagteg gecaaggagg tegagaceaa ggageagatt
                                                                       240
tetqccaecq eggeqatete egeeggegae acceagateg gegageteat egeegaggee
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Gln Ala Leu Val Arg Glu Gly Leu Arg Asn Val Ala Ala Gly Ala Asn
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Pro Leu Gly Leu Lys Arg Gly Ile Glu Lys Ala Val Glu Ala Val Thr
Gln Ser Leu Leu Lys Ser Ala Lys Glu Val Glu Thr Lys Glu Gln Ile
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Ser Ala Thr Ala Ala Ile Ser Ala Gly Asp Thr Gln Ile Gly Glu Leu
Ile Ala Glu Ala Met Asp Lys Val Gly Asn Glu Gly Val Ile Thr Val
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Glu Glu Ser Asn Thr Phe Gly Leu
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                            120
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# T	gcgctcacga tgagtgaccg catcgcggtg atgaacgccg gcaacgtcga acagatcggc	180 238
House St. St. and Market House Market House	agecegaceg agatetaega eegteeegeg aeggtgtteg tegeeagett categaat	238
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 	20 25 30	
	Tyr Val Thr His Asp Gln Glu Glu Ala Leu Thr Met Ser Asp Arg Ile	
	35 40 45	
	Ala Val Met Asn Ala Gly Asn Val Glu Gln Ile Gly Ser Pro Thr Glu 50 60	
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	cacatcaccg ctgcagaacc tgcagaacag acggcggatt ccgcggcacc gcccaagggc	360
	gcgccggtga tcgagatcga ccatgtcacg aagcgcttcg gcgactacct ggccgtcgcg	420
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245

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Val Ile Glu Ile Asp His Val Thr Lys Arg Phe Gly Asp Tyr Leu Ala

Gly	Ser	Thr	Leu 260	Lys	Ala	Arg	Pro	Gly 265	Glu.	Thr	Thr	Ile	Glu 270	Pro	Gly	
Gly	His	Ala 275		Leu	Met	Val	Arg 280		Glu	Arg	Ile	Arg 285		Thr	Pro	
Gly	Ser 290		Asp	Ala	Pro	Thr 295		Asp	Val	Ala	Cys 300		Arg	Ala	Thr	
Val 305		Asp	Leu	Thr	Phe 310		Gly	Pro	Val	Val 315		Leu	Ser	Leu	Ala 320	
	Pro	Asp	Asp	Ser 325		Val	Ile	Ala	His 330		Gly	Pro	Glu	Gln 335		
Leu	Pro	Leu	Leu 340		Pro	Gly	Asp	Asp 345	Val	Tyr	Val	Ser	Trp 350		Pro	
Glu	Ala	Ser 355		Val	Leu	Pro	Gly 360		Asp	Ile	Pro	Thr 365		Glu	Asp	
Leu	Glu 370		Met	Leu	Asp	Asp 375						303				
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gaga				gatco	a qa	atcqa	accat	. qt	2							33
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	_				_		_		_		_	_	_	_	atccgc	300
				ctggg						,5		55	٠ - ٠		5	323
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		212>		_												
			Mira					_								

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Val	Gln 210	Asp	Gly	Leu	Gly	Met 215	Ile	Met	Leu	Ser	Gln 220	Gly	Asn	Ser	Pro	
		Pro	Thr	Thr			Ile	Gln	Gln			Asp	Leu	Val	_	
225 Glu	Gln	Asn	Asp	Ara	230 Glv	Ser	Asp	Pro	Ser	235 Leu	His	Ara	Gln	Arg	240 Leu	
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Arg	Arg	Arg	Pro 260	Gly	Arg	Arg	Asn	Ile 265	Ala	Ile	Ala	Gln	Ala 270	Tyr	Ser	
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Val	Pro 290	Glu	Ser	Gly	Gly	Asp 295	Trp	Phe	Val	Asp	Thr 300	Met	Val	Ile	Pro	
Tyr 305		Thr	Gln	Asn	Gln 310		Ala	Ala	Glu	Ala 315		Ile	Asp	Tyr	Ile 320	
	Asp	Arg	Ala	Asn		Ala	Lys	Leu	Val		Phe	Thr	Gln	Phe		
D	21-	T	0	325	3 4 - 4	m\	3	a 1	330	37.	T	77- 7	3	335	3.7 -	
Pro	Ala	Leu	340	Asp	Met	Thr	Asp	345	Leu	Ala	Lys	Val	Asp 350	Pro	Ala	
Ser	Ala	Glu 355		Pro	Leu	Ile	Asn 360		Ser	Ala	Glu	Val 365		Ala	Asn	
Leu	Lys 370		Trp	Ala	Ala	Leu 375		Asp	Glu	Gln	Thr 380		Glu	Phe	Asn	
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-		210>														
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Tyr	Val	Asp 35	Ala	Phe	Thr	Met	Tyr 40	His	Glu	Gln	Ile	Phe 45	Arg	Ser	Phe	
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Ile	Leu	Pro	Met	165 Ala	Met	Pro	Gly	Val	170 Leu	Ala	Gly	Ser	Met	175 Leu	Val	
Phe	Ile	Pro	180 Ala	Val	Glv	Asp	Phe	185 Ile	Asn	Ala	Asp	Tyr	190 Leu	Gly	Ser	
		195			_	_	200				_	205		Leu		
	210					215					220			Met		
225	-	-	-		230					235		-			240	
			IIe	G1y 245	vai	Leu	ьeu	Tyr	250	Arg	Ата	ьеи	GIY	Ser 255	Glu	
Asp	Leu	Val														
		210>														
		211> 212>														
				bact	teri	ım va	accae	9								
ates		100>		accc.	~~ +-	a.c.c.c.	~~~+·	a	יבבבי	rtce	taa	1t t C	act (caacc	gacttc	6
															ccggc	12

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Glu Tyr Val Ile Met Gly Gln Val Leu Ser Ala Gly Ala Gly Gln Met
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Pro Ala Arg Gln Ala Ala Val Ala Ala Gly Ile Pro Trp Asp Val Ala
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LOOSISFI CIISOE
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ggtteggegg egetgaeget ggegatetae caecegeage agttecagta egeegggteg
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ctgtcgggct acctgaaccc gtccgagggg t ggtgacgcgg gcggctacaa ggccaacgac a gcctggaagc gcaacgaccc gatggtcaac a ctctc	atgtggggtc gcaccgagga cccgagcagc 420											
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gtgacgttcg cctccgacaa actcggcacg a												
ttcageggte agtacacett cageaegtee t gaeggeeegg egeegtegaa eeegaegatt o	-3-3-333											
aggcagtggg tgttcaacta caactggcag t	5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5											
cgcgagtacg ccgccgcgcg ttcgctggtg t												
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Asp Trp Ser Gly Arg Tyr Thr Val Va	al Thr Phe Ala Ser Asp Lys Leu											
35 40	45											
Gly Thr Ser Val Ala Ala Arg Gln Pr 50 55	co Glu Pro Asp Phe Ser Gly Gin 60											
Tyr Thr Phe Ser Thr Ser Cys Val Gl												
65 70	75 80											
Asp Gly Pro Ala Pro Ser Asn Pro Th												
85	90 95											
Thr Trp Asp Gly Arg Gln Trp Val Ph	ie wan the wan teh our teh our											

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100
                                105
Cys Phe Arq Gly Ala Asp Val Pro Arq Glu Tyr Ala Ala Ala Arg Ser
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Leu Val Phe Tyr Ala Pro Thr Ala Asp Gly Ser Met Phe Gly Thr Trp
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Arg Thr Asp Ile Leu Asp Gly Leu Cys Lys Gly Thr Val Ile Met Pro
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Val Ala Ala Tyr Pro Ala
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Gln Thr Ala Tyr Asp Thr Phe Met Ala Gly
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                                                                       300
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                                                                       360
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                                                                       420
cccqatqccq cccqcqtcq ccqaccacqt qqccqccqtt qtqqtcttcq gaaatccgtt
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                                    10
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                            40
Glu Pro Gly Leu Gly Trp Val Gly Asp Ala Phe Val Asn Ala Leu Arg
                        55
                                            60
Pro Lys Val Gly Glu Gln Ser Val Gly Thr Tyr Ala Val Asn Tyr Pro
                    70
                                        75
Ala Gly Phe Asp Phe Asp Lys Ser Ala Pro Met Gly Ala Ala Asp Ala
                                    90
Ser Gly Arg Val Gln Trp Met Ala Asp Asn Cys Pro Asp Thr Lys Leu
            100
                                105
Val Leu Gly Gly Met Ser Xaa Gly Ala Gly Val Ile Asp Leu Ile Thr
                            120
                                                 125
Val Asp Pro Arg Pro Leu Gly Arg Phe Thr Pro Thr Pro Met Pro Pro
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 Val
 Val
 Thr
 Lys
 Asp
 Glu
 Thr
 Thr
 Ile
 Val
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 Gly
 Ser
 Gly
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Glu Ala Thr Gly Ala Asn Ile Val Arq Val Ala Leu Ser Ala Pro Leu
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                                                 445
Lys Gln Ile Ala Phe Asn Gly Gly Leu Glu Pro Gly Val Val Ala Glu
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                                             460
Lys Val Ser Asn Leu Pro Ala Gly His Gly Leu Asn Ala Ala Thr Gly
465
                    470
                                         475
Glu Tyr Glu Asp Leu Leu Lys Ala Gly Val Ala Asp Pro Val Lys Val
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Thr Arg Ser Ala Leu Gln Asn Ala Ala Ser Ile Ala Ala Leu Phe Leu
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Thr Thr Glu Ala Val Val Ala Asp Lys Pro Glu
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                                                                       180
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                                                                       360
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                                                                       420
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                                                                       480
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                                                                       540
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                                                                       240
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                                                                       480
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                                                                       900
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Arg Gly Leu Asn Ala Leu Ala Asp Ala Val Lys Val Thr Leu Gly Pro
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Lys Gly Arg Asn Val Val Leu Glu Lys Lys Trp Gly Ala Pro Thr Ile
                            40
Thr Asn Asp Gly Val Ser Ile Ala Lys Glu Ile Glu Leu Glu Asp Pro
Tyr Glu Lys Ile Gly Ala Glu Leu Val Lys Glu Val Ala Lys Lys Thr
                    70
                                        75
Asp Asp Val Ala Gly Asp Gly Thr Thr Thr Ala Thr Val Leu Ala Gln
                                    90
Ala Leu Val Arg Glu Gly Leu Arg Asn Val Ala Ala Gly Ala Asn Pro
                                105
Leu Gly Leu Lys Arg Gly Ile Glu Lys Ala Val Glu Ala Val Thr Gln
       115
                            120
                                               125
Ser Leu Leu Lys Ser Ala Lys Glu Val Glu Thr Lys Glu Gln Ile Ser
                        135
Ala Thr Ala Ala Ile Ser Ala Gly Asp Thr Gln Ile Gly Glu Leu Ile
                                        155
Ala Glu Ala Met Asp Lys Val Gly Asn Glu Gly Val Ile Thr Val Glu
             165
                                    170
Glu Ser Asn Thr Phe Gly Leu Gln Leu Glu Leu Thr Glu Gly Met Arg
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Phe Asp Lys Gly Tyr Ile Ser Gly Tyr Phe Val Thr Asp Ala Glu Arg
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Gln Glu Ala Val Leu Glu Asp
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                            40
Asn Lys Ile Arg Gly Thr Phe Lys Ser Val Ala Val Lys Ala Pro Gly
                        55
                                            60
Phe Gly Asp Arg Arg Lys Ala Met Leu Gln Asp Met Ala Ile Leu Thr
                    70
                                        75
Gly Gly Gln Val Val Ser Glu Arg Val Gly Leu Ser Leu Glu Thr Ala
                                    90
Asp Val Ser Leu Leu Gly Gln Ala Arg Lys Val Val Thr Lys Asp
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. 110

105

Glu Thr Thr Ile Val Glu Gly Ser Gly Asp Ser Asp Ala Ile Ala Gly
115 120 125

Arg	Val 130	Ala	Gln	Ile	Arg	Ala 135	Glu	Ile	Glu	Asn	Ser 140	Asp	Ser	Asp	Tyr	
	Arg	Glu	Lys	Leu		Glu	Arg	Leu	Ala	-	Leu	Ala	Gly	Gly		
145	_	_		_	150	_	_			155	_				160	
Ala	Val	Ile	Lys	Ala 165	Gly	Ala	Ala	Thr	Glu 170	Val	Glu	Leu	Lys	Glu 175	Arg	
Lys	His	Arg	Ile 180	Glu	Asp	Ala	Val	Arg 185	Asn	Ala	Lys	Ala	Ala 190	Val	Glu	
Glu	Gly	Ile 195	Val	Ala	Gly	Gly	Gly. 200	Val	Ala	Leu	Leu	Gln 205	Ser	Ala	Pro	
Ala	Leu 210	Asp	Asp	Leu	Gly	Leu 215	Thr	Gly	Asp	Glu	Ala 220	Thr	Gly	Ala	Asn	
Ile 225		Arg	Val	Ala	Leu 230		Ala	Pro	Leu	Lys 235		Ile	Ala	Phe	Asn 240	
	Gly	Leu	Glu	Pro 245		Val	Val	Ala	Glu 250		Val	Ser	Asn	Leu 255		
Ala	Gly	His			Asn	Ala	Ala			Glu	Tyr	Glu	-	Leu	Leu	
Lys	Ala	-	260 Val	Ala	Asp	Pro		265 Lys	Val	Thr	Arg		270 Ala	Leu	Gln	
Asn		275 Ala	Ser	Ile	Ala		280 Leu	Phe	Leu	Thr		285 Glu	Ala	Val	Val	
Ala 305	290 Asp	Lys	Pro	Glu		295					300					
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	<2		unst (973		. (973	3)										
gato				gaad	et co	acct	gatt	. aa	cctac	acc	atco	caat	ca o	caato	cgggtt	60
									-		-	_	_		cagcgc	120
															gctgct	180
															gccac	240
															catcca	300
ggg	cgcad	cca ç	gaaga	acago	et go	gegea	iggcg	g gat	tccg	gtcg	atct	tcct	.cg a	acgto	cgcgcg	360
ctto	cgcg	ctg a	atcgo	ggto	g gt	atca	ccgt	: gat	cate	gcc	tato	gtcto	ggg g	gcgcg	gaacgt	420
gggg	gggc	ctg t	tcac	cgca	ac to	ggcc	gtcac	tto	cato	gtt	cttg	gcct	.gg (ctctg	gcagaa	480
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cgactggate accgtececa ecgeggeggg ecggeegtee geecaeggee gegtggtgga
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                                25
Ala Leu Arg Arg Arg Gly Ser Ala Leu Ala Arg Pro Val Gln Leu Leu
                            40
                                                45
Arg Thr Tyr Ile Leu Pro Leu Gly Ala Leu Leu Leu Leu Val Gln
Ala Met Glu Ile Ser Asp Asp Ala Thr Ser Val Arg Leu Val Ala Thr
                                        75
Leu Phe Gly Val Val Leu Leu Thr Leu Val Leu Ser Gly Leu Asn Ala
                                    90
Thr Leu Ile Gln Gly Ala Pro Glu Asp Ser Trp Arg Arg Ile Pro
            100
                                105
                                                    110
Ser Ile Phe Leu Asp Val Ala Arg Phe Ala Leu Ile Ala Val Gly Ile
                            120
Thr Val Ile Met Ala Tyr Val Trp Gly Ala Asn Val Gly Gly Leu Phe
                        135
                                            140
Thr Ala Leu Gly Val Thr Ser Ile Val Leu Gly Leu Ala Leu Gln Asn
                    150
                                        155
Ser Val Gly Gln Ile Ile Ser Gly Leu Leu Leu Phe Glu Gln Pro
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Phe Arg Leu Gly Asp Trp Ile Thr Val Pro Thr Ala Ala Gly Arg Pro
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                                185
Ser Ala His Gly Arg Val Val Glu Val Asn Trp Arg Ala Thr His Ile
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600

660

720

780

840

900

960

1020

1080

1140

1200

1260

1320

1366

205

200

215

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<212> DNA

<213> Mycobacterium vaccae

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                                                             240
Val Val Thr Thr Phe Asn Ala Ala Asp Thr Pro Asp Asp Val Cys Glu
                                    250
Met Leu Ser Ser Val Ala Ala Ser Leu Pro Glu Leu Arg Thr Asp Gly
                                265
Gln Ile Ala Thr Leu Tyr Leu Gly Ala Ala Glu Tyr Glu Lys Ser Ile
                            280
Pro Leu His Thr Pro Ala Val Asp Asp Ser Val Arg Ser Thr Tyr Leu
                        295
                                            300
Arg Trp Val Trp Tyr Ala Ala Arg Arg Gln Glu Leu Arg Xaa Asn Gly
                                        315
Val Ala Asp Xaa Phe Asp Thr Pro Glu Arg Ile Ala Ser Ala Met Arg
                325
                                    330
Ala Val Ala Ser Thr Leu Arg Leu Ala Asp Asp Glu Gln Gln Glu Ile
                                345
Ala Asp Val Val Arg Leu Val Arg Tyr Gly Asn Gly Glu Arg Leu Gln
                            360
Gln Pro Gly Gln Val Pro Thr Gly Met Arg Phe Ile Val Asp Gly Arg
                        375
                                            380
Val Ser Leu Ser Val Ile Asp Gln Asp Gly Asp Val Ile Pro Ala Arg
                    390
                                        395
Val Leu Glu Arg Gly Asp Phe Leu Gly Gln Thr Thr Leu Thr Arg Glu
                405
                                    410
Pro Val Leu Ala Thr Ala His Ala Leu Glu Glu Val Thr Val Leu Glu
            420
                                425
Met Ala Arg Asp Glu Ile Glu Arg Leu Val His Arg Lys Pro Ile Leu
                            440
Leu His Val Ile Gly Ala Val
    450
      <210> 122
      <211> 898
      <212> DNA
      <213> Mycobacterium vaccae
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egetaceace teetgtegeg gatgageate cagteeaagt tgetgetgat getgettetg
                                                                       120
accagcatte teteggetge ggtggteggt tteategget atcagteegg aeggteeteg
                                                                       180
etgegegeat eggtgttega eegecteaec gacateegeg agtegeagte gegegggttg
                                                                       240
gagaatcagt tegeggaeet gaagaacteg atggtgattt actegegegg cageactgee
                                                                       300
                                                                       360
acqqaqqcqa tcqqcqcqtt caqcqacqgt ttccqtcagc tcqqcgatgc gacqatcaat
accqqqcaqq cqqcqtcatt qcqccqttac tacqaccqqa cqttcqccaa caccaccctc
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gacgacagcg gaaaccgcgt cgacgtccgc gcgctcatcc cgaaatccaa cccccagcgc
                                                                       480
tatctgcagg cgctctatac cccgccgttt cagaactggg agaaggcgat cgcgttcgac
                                                                       540
qacqcqcqcq acqqcaqcqc ctqqtcqqcc qccaatqcca qattcaacqa qttcttccgc
                                                                       600
gagategtge accgetteaa ettegaggat etgatgetge tegacetega gggeaacgtg
                                                                       660
gtgtactccg cctacaaggg gccggatctc gggacaaaca tcgtcaacgg cccctatcgc
                                                                       720
                                                                       780
aaccgggaac tgtcggaagc ctacgagaag gcggtcgcgt cgaactcgat cgactatgtc
qqtqtcaccq acttcggqtq gtacctgcct gccgaggaac cgaccgcctg gttcctgtcc
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                                                                       898
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      <211> 1259
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<210> 124

<211> 299

<212> PRT

<213> Mycobacterium vaccae

<400> 124

210

Met Thr Ile Leu Pro Trp Asn Ala Arg Thr Ser Glu His Pro Thr Arg 10 Lys Arg Arg Gly Arg Tyr His Leu Leu Ser Arg Met Ser Ile Gln Ser 20 25 Lys Leu Leu Met Leu Leu Thr Ser Ile Leu Ser Ala Ala Val 40 45 Val Gly Phe Ile Gly Tyr Gln Ser Gly Arg Ser Ser Leu Arg Ala Ser Val Phe Asp Arg Leu Thr Asp Ile Arg Glu Ser Gln Ser Arg Gly Leu 75 70 Glu Asn Gln Phe Ala Asp Leu Lys Asn Ser Met Val Ile Tyr Ser Arg 90 Gly Ser Thr Ala Thr Glu Ala Ile Gly Ala Phe Ser Asp Gly Phe Arg 105 Gln Leu Gly Asp Ala Thr Ile Asn Thr Gly Gln Ala Ala Ser Leu Arg 115 120 Arg Tyr Tyr Asp Arg Thr Phe Ala Asn Thr Thr Leu Asp Asp Ser Gly 135 140 Asn Arg Val Asp Val Arg Ala Leu Ile Pro Lys Ser Asn Pro Gln Arg 150 155 Tyr Leu Gln Ala Leu Tyr Thr Pro Pro Phe Gln Asn Trp Glu Lys Ala 165 170 Ile Ala Phe Asp Asp Ala Arg Asp Gly Ser Ala Trp Ser Ala Ala Asn 185 Ala Arg Phe Asn Glu Phe Phe Arg Glu Ile Val His Arg Phe Asn Phe 200 205 Glu Asp Leu Met Leu Leu Asp Leu Glu Gly Asn Val Val Tyr Ser Ala

215

<210> 125

<211> 419

<212> PRT

<213> Mycobacterium vaccae

<400> 125

Gln Leu Met Thr Ala Arg Gly Gln Trp Arg Asp Thr Gly Met Gly Asp Thr Gly Glu Thr Ile Leu Val Gly Pro Asp Asn Leu Met Arg Ser Asp Ser Arg Leu Phe Arg Glu Asn Arg Glu Lys Phe Leu Ala Asp Val Val 40 Glu Gly Gly Thr Pro Pro Glu Val Ala Asp Glu Ser Val Asp Arg Arg 55 Gly Thr Thr Leu Val Gln Pro Val Thr Thr Arg Ser Val Glu Glu Ala Gln Arg Gly Asn Thr Gly Thr Thr Ile Glu Asp Asp Tyr Leu Gly His 90 Glu Ala Leu Gln Ala Tyr Ser Pro Val Asp Leu Pro Gly Leu His Trp 105 Val Ile Val Ala Lys Ile Asp Thr Asp Glu Ala Phe Ala Pro Val Ala 120 Gln Phe Thr Arg Thr Leu Val Leu Ser Thr Val Ile Ile Phe Gly 135 140 Val Ser Leu Ala Ala Met Leu Leu Ala Arg Leu Phe Val Arg Pro Ile 150 155 Arg Arg Leu Gln Ala Gly Ala Gln Gln Ile Ser Gly Gly Asp Tyr Arg 165 170 Leu Ala Leu Pro Val Leu Ser Arg Asp Glu Phe Gly Asp Leu Thr Thr 185 Ala Phe Asn Asp Met Ser Arg Asn Leu Ser Ile Lys Asp Glu Leu Leu 200 Gly Glu Glu Arg Ala Glu Asn Gln Arg Leu Met Leu Ser Leu Met Pro 215 220 Glu Pro Val Met Gln Arg Tyr Leu Asp Gly Glu Glu Thr Ile Ala Gln 235 230 Asp His Lys Asn Val Thr Val Ile Phe Ala Asp Met Met Gly Leu Asp 245 250 Glu Leu Ser Arg Met Leu Thr Ser Glu Glu Leu Met Val Val Asn 265 Asp Leu Thr Arg Gln Phe Asp Ala Ala Ala Glu Ser Leu Gly Val Asp 280 His Val Arg Thr Leu His Asp Gly Tyr Leu Ala Ser Cys Gly Leu Gly 295 300 Val Pro Arg Leu Asp Asn Val Arg Arg Thr Val Asn Phe Ala Ile Glu

310

<400> 129

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Met Asp Arg Ile Ile Asp Arg His Ala Ala Glu Ser Gly His Asp Leu
Arg Leu Arg Ala Gly Ile Asp Thr Gly Ser Ala Ala Ser Gly Leu Val
                                345
Gly Arg Ser Thr Leu Ala Tyr Asp Met Trp Gly Ser Ala Val Asp Val
                            360
Ala Tyr Gln Val Gln Arg Gly Ser Pro Gln Pro Gly Ile Tyr Val Thr
                        375
Ser Arg Val His Glu Val Met Gln Glu Thr Leu Asp Phe Val Ala Ala
                    390
                                        395
Gly Glu Val Val Gly Glu Arg Gly Val Glu Thr Val Trp Arg Leu Gln
                405
                                    410
Gly His Pro
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      <211> 27
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <400> 126
ccggatccga tgagcagcgt gctgaac
                                                                         27
      <210> 127
      <211> 26
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <400> 127
                                                                         26
gcggatccca cggccccgat cacgtg
      <210> 128
      <211> 33
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <400> 128
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ccggatccaa tgacatttct gccctggaat gcg
      <210> 129
      <211> 32
      <212> DNA
      <213> Artificial Sequence
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	.000	
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	<2223> Made III a lab	
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	ccggatccgg agcaaccgtt ccggctc	27
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	<220> <223> Made in a lab	
	(225) Hade In a Tab	
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	<211> 844	
i	<212> DNA	
	<213> Mycobacterium vaccae	
	<400> 132	60
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	gcccacggcc gcgtggtgga agtcaactgg cgtgcaacac atatcgacac cggcggcaac	180
	ctgctggtaa tgcccaacgc cgaactcgcc ggcgcgtcgt tcaccaatta cagccggccc gtgggagagc accggctgac cgtcgtcacc accttcaacg ccgcggacac ccccgatgat	240
	gtotgogaga tgotgtogto ggtogoggog togotgocog aactgogoac ogacggacag	300
i	ategecacge tetatetegg tgeggeegaa tacgagaagt egateeegtt geacacacee	360
; !	gcggtggacg actcggtcag gagcacgtac ctgcgatggg tctggtacgc cgcgcgcgg	420
1	caggaacttc gcctaacggc gtcgccgacg attcgacacg ccggaacgga tcgcctcggc	480
į	catgogggot gtggcgtcca cactgogott ggcagacgac gaacagcagg agatogccga	540
	cgtggtgcgt ctggtccgtt acggcaacgg ggaacgcctc cagcagccgg gtcaggtacc	600
	gaccgggatg aggttcatcg tagacggcag ggtgagtctg tccgtgatcg atcaggacgg	660
	cgacgtgatc ccggcgcggg tgctcgagcg tggcgacttc ctggggcaga ccacgctgac	720
	gcgggaaccg gtactggcga ccgcgcacgc gctggaggaa gtcaccgtgc tggagatggc	780
	ccgtgacgag atcgagcgcc tggtgcaccg aaagccgatc ctgctgcacg tgatcggggc	840
	cgtg	844
	<210> 133	
	<211> 742	
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	<213> Mycobacterium vaccae	
	<400> 133	
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	atttactcgc gcggcagcac tgccacggag gcgatcggcg cgttcagcga cggtttccgt	180
	cageteggeg atgegaegat caataceggg caggeggegt cattgegeeg ttactacgae	240
	cggacgttcg ccaacaccac cctcgacgac agcggaaacc gcgtcgacgt ccgcgcgctc	300

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atcccqaaat ccaacccca qcqctatctq caqqcqctct ataccccqcc gtttcagaac
tgggagaagg cgatcgcgtt cgacgacgcg cgcgacggca gcgcctggtc ggccgccaat
gccagattca acgagttctt ccgcgagatc gtgcaccgct tcaacttcga ggatctgatg
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aacatcgtca acggccccta tcgcaaccgg gaactgtcgg aagcctacga gaaggcggtc
gegtegaact egategaeta tgteggtgte accgaetteg ggtggtaeet geetgeegag
gaaccgaccg cctggttcct gtccccggtc gggttgaagg accgagtcga cggtgtgatg
gcggtccagt tccccggaat tc
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      <211> 282
      <212> PRT
      <213> Mycobacterium vaccae
      <220>
      <221> UNSURE
      <222> (145)...(145)
      <221> UNSURE
      <222> (151)...(151)
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Gly Arg Pro Ser Ala His Gly Arg Val Val Glu Val Asn Trp Arg Ala
Thr His Ile Asp Thr Gly Gly Asn Leu Leu Val Met Pro Asn Ala Glu
                            40
                                                 45
Leu Ala Gly Ala Ser Phe Thr Asn Tyr Ser Arg Pro Val Gly Glu His
Arg Leu Thr Val Val Thr Thr Phe Asn Ala Ala Asp Thr Pro Asp Asp
                    70
                                        75
Val Cys Glu Met Leu Ser Ser Val Ala Ala Ser Leu Pro Glu Leu Arg
                85
                                    90
Thr Asp Gly Gln Ile Ala Thr Leu Tyr Leu Gly Ala Ala Glu Tyr Glu
            100
                                105
Lys Ser Ile Pro Leu His Thr Pro Ala Val Asp Asp Ser Val Arg Ser
                            120
Thr Tyr Leu Arg Trp Val Trp Tyr Ala Ala Arg Arg Gln Glu Leu Arg
                        135
Xaa Asn Gly Val Ala Asp Xaa Phe Asp Thr Pro Glu Arg Ile Ala Ser
                    150
                                        155
Ala Met Arg Ala Val Ala Ser Thr Leu Arg Leu Ala Asp Asp Glu Gln
                                    170
Gln Glu Ile Ala Asp Val Val Arg Leu Val Arg Tyr Gly Asn Gly Glu
                                185
Arg Leu Gln Gln Pro Gly Gln Val Pro Thr Gly Met Arg Phe Ile Val
        195
                            200
                                                 205
Asp Gly Arg Val Ser Leu Ser Val Ile Asp Gln Asp Gly Asp Val Ile
                        215
Pro Ala Arg Val Leu Glu Arg Gly Asp Phe Leu Gly Gln Thr Thr Leu
                    230
                                        235
Thr Arg Glu Pro Val Leu Ala Thr Ala His Ala Leu Glu Glu Val Thr
                                    250
Val Leu Glu Met Ala Arg Asp Glu Ile Glu Arg Leu Val His Arg Lys
                                265
```

420

480

540

600

660

720

742

Pro Ile Leu Leu His Val Ile Gly Ala Val

275 280 <210> 135 <211> 247 <212> PRT <213> Mycobacterium vaccae <400> 135 Gly Tyr Gln Ser Gly Arg Ser Ser Leu Arg Ala Ser Val Phe Asp Arg 10 Leu Thr Asp Ile Arg Glu Ser Gln Ser Arg Gly Leu Glu Asn Gln Phe 25 Ala Asp Leu Lys Asn Ser Met Val Ile Tyr Ser Arg Gly Ser Thr Ala Thr Glu Ala Ile Gly Ala Phe Ser Asp Gly Phe Arg Gln Leu Gly Asp 55 Ala Thr Ile Asn Thr Gly Gln Ala Ala Ser Leu Arg Arg Tyr Tyr Asp 70 Arg Thr Phe Ala Asn Thr Thr Leu Asp Asp Ser Gly Asn Arg Val Asp 90 Val Arg Ala Leu Ile Pro Lys Ser Asn Pro Gln Arg Tyr Leu Gln Ala 100 105 Leu Tyr Thr Pro Pro Phe Gln Asn Trp Glu Lys Ala Ile Ala Phe Asp 120 125 Asp Ala Arg Asp Gly Ser Ala Trp Ser Ala Ala Asn Ala Arg Phe Asn Glu Phe Phe Arg Glu Ile Val His Arg Phe Asn Phe Glu Asp Leu Met 150 155 Leu Leu Asp Leu Glu Gly Asn Val Val Tyr Ser Ala Tyr Lys Gly Pro 170 Asp Leu Gly Thr Asn Ile Val Asn Gly Pro Tyr Arg Asn Arg Glu Leu 185 Ser Glu Ala Tyr Glu Lys Ala Val Ala Ser Asn Ser Ile Asp Tyr Val 200 Gly Val Thr Asp Phe Gly Trp Tyr Leu Pro Ala Glu Glu Pro Thr Ala 215 220 Trp Phe Leu Ser Pro Val Gly Leu Lys Asp Arg Val Asp Gly Val Met 230 235 Ala Val Gln Phe Pro Gly Ile 245 <210> 136 <211> 45 <212> DNA <213> Mycobacterium vaccae <220> <221> unsure <222> (18) ... (18) <400> 136 atgagcgaaa tcgcccgncc ctggcgggtt ctggcatgtg gcatc

45

<210> 137

<211> 340

<212> DNA

<213> Mycobacterium vaccae

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      <222> (273)...(273)
      <221> unsure
      <222> (286)...(286)
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                                                                         60
cccgcgatge ccgcccgccc ggtgtccacg atcgcgccgg cgacctcggg cacgctcagc
                                                                        120
gagtttttcg ccgccaaggg cgtcacgatg gagccgcagt ccagccgcga cttccgcgcc
                                                                        180
ctcaacatcg tgctgccgaa gccgcggggc tgggagcaca tcccggaccc gaacgtgccg
                                                                        240
gacgcgttcg cggtgctggc cgaccgggtc agnggtaaag gtcagnagtc gacaaacgcc
                                                                        300
cacgtggtgg tcgacaaaca cgtaggcgag ttcgacggca
                                                                        340
      <210> 138
      <211> 235
      <212> DNA
      <213> Mycobacterium vaccae
      <220>
      <221> unsure
      <222> (16)...(16)
      <400> 138
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                                                                         60
caacggette aaggteageg tteegggtee gggteeggee geacegeeae etgeaeeegg
                                                                        120
tgcccccggt gtcccgcccg cccccggcgc cccggcgctg ccgctggccg tcgcaccacc
                                                                        180
cccggctccc gctgttcccg ccgtggcgcc cgcgccacag ctgctgggac tgcag
                                                                        235
      <210> 139
      <211> 15
      <212> PRT
      <213> Mycobacterium vaccae
      <400> 139
Met Ser Glu Ile Ala Arg Pro Trp Arg Val Leu Ala Cys Gly Ile
 1
                 5
                                     10
      <210> 140
      <211> 113
      <212> PRT
      <213> Mycobacterium vaccae
      <220>
      <221> UNSURE
      <222> (96) ... (96)
      <400> 140
Ala Thr Gly Gly Ala Ala Ala Val Pro Ala Gly Val Ser Ala Pro Ala
                 5
 1
Val Ala Pro Ala Pro Ala Met Pro Ala Arg Pro Val Ser Thr Ile Ala
Pro Ala Thr Ser Gly Thr Leu Ser Glu Phe Phe Ala Ala Lys Gly Val
                            40
Thr Met Glu Pro Gln Ser Ser Arg Asp Phe Arg Ala Leu Asn Ile Val
```

```
50
                                            60
                        55
Leu Pro Lys Pro Arg Gly Trp Glu His Ile Pro Asp Pro Asn Val Pro
                    70
                                        75
Asp Ala Phe Ala Val Leu Ala Asp Arg Val Gly Lys Gly Gln Xaa
                                    90
Ser Thr Asn Ala His Val Val Val Asp Lys His Val Gly Glu Phe Asp
                                105
Gly
      <210> 141
      <211> 73
      <212> PRT
      <213> Mycobacterium vaccae
      <400> 141
Val Thr Thr Ser Val Glu Gln Val Val Ala Ala Asp Ala Thr Glu
Ala Ile Val Asn Gly Phe Lys Val Ser Val Pro Gly Pro Gly Pro Ala
                                25
Ala Pro Pro Pro Ala Pro Gly Ala Pro Gly Val Pro Pro Ala Pro Gly
                            40
                                                45
Ala Pro Ala Leu Pro Leu Ala Val Ala Pro Pro Pro Ala Pro Ala Val
                        55
Pro Ala Val Ala Pro Ala Pro Gln Leu
      <210> 142
      <211> 273
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 142
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                                                                        60
agcaacgccg cggccaaggg ctacttcccg ctgagcttca ccgtcgccgg catcgaccag
                                                                       120
aacggtccga tcgtgaccgc caacgtcacc gcggcggccc cgacgggcgc cgtggccacc
                                                                       180
caqccqctqa cqttcatcqc cqqqccqaqc ccqaccggat ggcagctqtc caagcagtcc
                                                                       240
gcactggccc tgatgtccgc ggtcatcgcc gca
                                                                       273
      <210> 143
      <211> 91
      <212> PRT
      <213> Mycobacterium vaccae
      <400> 143
Ala Thr Tyr Val Gln Gly Gly Leu Gly Arg Ile Glu Ala Arg Val Ala
Asp Ser Gly Tyr Ser Asn Ala Ala Ala Lys Gly Tyr Phe Pro Leu Ser
                                25
Phe Thr Val Ala Gly Ile Asp Gln Asn Gly Pro Ile Val Thr Ala Asn
Val Thr Ala Ala Ala Pro Thr Gly Ala Val Ala Thr Gln Pro Leu Thr
Phe Ile Ala Gly Pro Ser Pro Thr Gly Trp Gln Leu Ser Lys Gln Ser
                    70
Ala Leu Ala Leu Met Ser Ala Val Ile Ala Ala
                85
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<210> 144
      <211> 554
      <212> DNA
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                                                                        60
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caagattcga aggacccaaa caacatgaaa ttcactggaa tgaccgtgcg cgcaagccgc
                                                                        180
gegecetgge eggegteggg geggeatgte tgtteggegg egtggeegeg geaacegtgg
                                                                        240
eggeacagat ggegggegec cageeggeeg agtgeaaege cageteaete aeeggeaeeg
                                                                        300
teageteggt gaceggteag gegegteagt acetagaeae eeaceeggge gecaaceagg
                                                                        360
ccgtcaccgc ggcgatgaac cagccgcqcc ccgaqqccqa qqcgaacctq cqqqqctact
                                                                        420
teacegecaa eceggeggag tactaegace tgeggggeat ectegeeceg ateggtgaeg
                                                                        480
cgcagcgcaa ctgcaacatc accgtgctgc cggtagagct gcagacggcc tacgacacgt
                                                                        540
tcatqqccqq ctqa
                                                                        554
      <210> 145
      <211> 136
      <212> PRT
      <213> Mycobacterium vaccae
      <400> 145
Met Lys Phe Thr Gly Met Thr Val Arg Ala Ser Arg Arg Ala Leu Ala
                 5
Gly Val Gly Ala Ala Cys Leu Phe Gly Gly Val Ala Ala Ala Thr Val
Ala Ala Gln Met Ala Gly Ala Gln Pro Ala Glu Cys Asn Ala Ser Ser
                            40
Leu Thr Gly Thr Val Ser Ser Val Thr Gly Gln Ala Arq Gln Tyr Leu
                        55
Asp Thr His Pro Gly Ala Asn Gln Ala Val Thr Ala Ala Met Asn Gln
                    70
                                         75
Pro Arg Pro Glu Ala Glu Ala Asn Leu Arg Gly Tyr Phe Thr Ala Asn
                85
                                     90
Pro Ala Glu Tyr Tyr Asp Leu Arg Gly Ile Leu Ala Pro Ile Gly Asp
                                 105
Ala Gln Arg Asn Cys Asn Ile Thr Val Leu Pro Val Glu Leu Gln Thr
                            120
Ala Tyr Asp Thr Phe Met Ala Gly
    130
                        135
      <210> 146
      <211> 808
      <212> DNA
      <213> Mycobacterium vaccae
      <220>
      <221> unsure
      <222> (15)...(15)
      <400> 146
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                                                                        60
gaatcacccg tgtgccaatt cagtgcggc aacggtgtcc gtccacgaag ggattcagga
                                                                       120
aatgatgaca actcgccgga agtcagccgc agtggcggga atcgctgcgg tggccatcct
                                                                       180
eggtgeggee geatgttega gtgaggaegg tgggageaeg geetegtegg eeageageae
                                                                       240
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<221> unsure

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ggccgaccct gcggccaacc tgatcggctc cggctgcgcg gcctacgccg agcaggtccc
cgaaggtccc gggtcggtgg ccgggatggc agccgatccg gtgacggtgg cggcgtcgaa
caacceqatg ctgcagacgc tgtcccaggc gctgtccggc cagctcaatc cgcaggtcaa
tetegtegae accetegaeg geggtgagtt cacegtgtte gegeegaeeg acgaegegtt
cgccaagatc gatccggcca cgctggagac cctcaagacg gactccgaca tgctgaccaa
catectgace taccaegteg tgeeeggeea ggeegegeee gateaggtgg teggegagea
tgtgacggtg gagggggcgc cggtcacggt gtccgggatg gccgaccagc tcaaggtcaa
cgacgcgtcg gtggtgtgcg gtggggtgca gaccgccaac gcgacggtgt atctgatcga
caccgtgctg atgccgccgg cagcgtag
      <210> 147
      <211> 228
      <212> PRT
      <213> Mycobacterium vaccae
      <400> 147
Met Met Thr Thr Arg Arg Lys Ser Ala Ala Val Ala Gly Ile Ala Ala
Val Ala Ile Leu Gly Ala Ala Ala Cys Ser Ser Glu Asp Gly Gly Ser
Thr Ala Ser Ser Ala Ser Ser Thr Ala Ser Ser Ala Met Glu Ser Ala
                            40
Thr Asp Glu Met Thr Thr Ser Ser Ala Ala Pro Ser Ala Asp Pro Ala
Ala Asn Leu Ile Gly Ser Gly Cys Ala Ala Tyr Ala Glu Gln Val Pro
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Glu Gly Pro Gly Ser Val Ala Gly Met Ala Ala Asp Pro Val Thr Val
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Ala Ser Asn Asn Pro Met Leu Gln Thr Leu Ser Gln Ala Leu Ser
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Gly Gln Leu Asn Pro Gln Val Asn Leu Val Asp Thr Leu Asp Gly Gly
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Glu Phe Thr Val Phe Ala Pro Thr Asp Asp Ala Phe Ala Lys Ile Asp
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                                            140
Pro Ala Thr Leu Glu Thr Leu Lys Thr Asp Ser Asp Met Leu Thr Asn
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Ile Leu Thr Tyr His Val Val Pro Gly Gln Ala Ala Pro Asp Gln Val
                                    170
                                                         175
                165
Val Gly Glu His Val Thr Val Glu Gly Ala Pro Val Thr Val Ser Gly
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                                185
Met Ala Asp Gln Leu Lys Val Asn Asp Ala Ser Val Val Cys Gly Gly
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	Leu 145		Thr	Lys	Ile	Leu 150		Tyr	His	Val	Val 155		Gly	Gln	Ala	Ala 160		
4		Asp	Gln	Val	Val 165		Glu	His	Val	Thr 170		Glu	Gly	Ala	Pro 175			
L	Thr	Val	Ser	_		Ala	Asp	Gln	Leu 185		Val	Asn	Asp	Ala 190	Ser	Val		
- -	Val	Cys	Gly 195	180 Gly	Val	Gln	Thr	Ala 200		Ala	Thr	Val	Tyr 205		Ile	Asp		
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cccgatgccg ccccgcgtcg ccgaccacgt ggccgccgtt gtggtcttcg gaaatccgtt
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                                                                      1080
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Ser Gly Arg Val Gln Trp Met Ala Asp Asn Cys Pro Asp Thr Lys Leu 105 110 100 Val Leu Gly Gly Met Ser Xaa Gly Ala Gly Val Ile Asp Leu Ile Thr

120 125 115

Val Asp Pro Arg Pro Leu Gly Arg Phe Thr Pro Thr Pro Met Pro Pro 135 140

Arq Val Ala Asp His Val Ala Ala Val Val Phe Gly Asn Pro Leu 150 155

Arg Asp Ile Arg Gly Gly Pro Arg Leu Glu Pro Arg Gly Leu Asn 170

Met Glu Thr Ser Glu Arg Gly Leu Tyr Thr His Arg Thr Tyr Arg Gly 180 185 190

Leu Tyr Pro Arg Leu Tyr Ser Ser Glu Arg Ile Leu Glu Ala Ser Pro 200

Leu Glu Cys Tyr Ser Ala Leu Ala Leu Glu Ala Ser Pro Ala Ser Pro 220 215

Pro Arq Pro His Glu Cys Tyr Ser Ser Glu Arg Pro Arg Gly Leu Tyr 230 235

Pro His Glu Ala Ser Asn Leu Glu Pro Arg Ala Leu Ala His Ile Ser 245 250

Pro His Glu Ala Leu Ala Thr Tyr Arq Ala Leu Ala Ala Ser Pro Ala 265 Ser Asn Gly Leu Tyr Met Glu Thr Val Ala Leu Gly Leu Gly Leu Ala 280 Leu Ala Ala Leu Ala Ala Ser Asn Pro His Glu Ala Leu Ala Ala Arq 295 Gly Leu Glu Gly Leu Pro Arg Gly Leu Tyr Gly Leu Asn Ser Glu Arg 310 315 Val Ala Leu Gly Leu Leu Glu Pro Arg Gly Leu Ala Leu Ala Pro Arg 325 330 Thr Tyr Arg Leu Glu His Ile Ser Leu Glu Pro His Glu Val Ala Leu 345 Pro Arg Ala Arg Gly Gly Leu Tyr Gly Leu Val Ala Leu Thr His Arg 360 Leu Glu Gly Leu Ala Ser Pro Ala Leu Ala Gly Leu Tyr Pro Arg Leu 375 380 Glu Ala Arg Gly Gly Leu Gly Leu Tyr Ala Ser Pro Ala Leu Ala Val 390 395 Ala Leu Ala Arg Gly Pro His Glu Thr His Arg Ala Leu Ala Ser Glu 405 410 Arg Gly Leu Tyr Gly Leu Tyr Gly Leu Asn Ala Arg Gly Val Ala Leu 420 425 Thr His Arg Ala Leu Ala Thr His Arg Al'a Leu Ala Pro Arg Ala Leu 440 Ala Gly Leu Ile Leu Glu Leu Glu Val Ala Leu Thr Arg Pro Gly Leu 455 Met Glu Thr His Ile Ser Ala Leu Ala Gly Leu Tyr Leu Glu Gly Leu 470 475 Tyr Ala Leu Ala Ala Leu Ala Ala Leu Ala Ala Ser Asn Ala Arg Gly 490 Ser Glu Arg Pro Arg Ala Leu Ala Gly Leu Tyr Ala Arg Gly Ala Arg 500 505 Gly Ser Glu Arg Thr His Arg Ala Leu Ala Ala Arg Gly Ala Arg Gly 520 Met Glu Thr His Ile Ser Ile Leu Glu Ala Arg Gly Thr His Arg Thr 535 540 Arg Pro Thr His Arg Ala Arg Gly Ala Leu Ala Val Ala Leu Gly Leu 550 555 Tyr Gly Leu Tyr Thr His Arg Gly Leu Tyr Pro Arg Ala Ser Pro Gly 570 Leu His Ile Ser Ala Ser Asn Ala Arg Gly Gly Leu Ala Arg Gly Thr 585 His Arg Ala Arg Gly Ser Glu Arg Ala Arg Gly Thr His Arg Ala Arg 600 Gly Pro Arg Gly Leu Tyr Ala Ser Pro Ala Arg Gly Ala Arg Gly Gly 615 620 Leu Tyr Ala Arg Gly Ala Arg Gly Thr His Arg Ser Glu Arg Gly Leu 630 635 Tyr Ala Arg Gly Ala Arg Gly Ala Leu Ala Val Ala Leu Ala Arg Gly 650 Pro Arg Ala Leu Ala Pro Arg Gly Leu Tyr Gly Leu Tyr Pro Arg Gly 665 Leu Tyr Ala Leu Ala Ala Leu Ala Gly Leu Tyr Leu Glu Ala Ser Pro Ala Leu Ala Val Ala Leu Gly Leu Tyr Val Ala Leu Gly Leu Tyr Ala 695 Leu Ala Ala Ser Pro Ala Arg Gly Pro Arg Ala Leu Ala Ala Leu Ala

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His Leu Thr Ile Gln Gln Trp Asp Thr Phe Leu Asn Gly Val Phe Pro
Leu Asp Arg Asn Arg Leu Thr Arg Glu Trp Phe His Ser Gly Lys Ala
                    70
                                         75
Thr Tyr Val Val Ala Gly Glu Gly Ala Asp Glu Phe Glu Gly Thr Leu
                                     90
Glu Leu Gly Tyr His Val Gly Phe Pro Trp Ser Leu Gly Val Gly Ile
                                105
Asn Phe Ser Tyr Thr Thr Pro Asn Ile Thr Tyr Asp Gly Tyr Gly Leu
                            120
Asn Phe Ala Asp Pro Leu Leu Gly Phe Gly Asp Ser Ile Val Thr Pro
                        135
                                            140
Pro Leu Phe Pro Gly Val Ser Ile Thr Ala Asp Leu Gly Asn Gly Pro
                    150
                                        155
Gly Ile Gln Glu Val Ala Thr Phe Ser Val Asp Val Ala Gly Pro Gly
                                     170
                165
Gly Ser Val Val Val Ser Asn Ala His Gly Thr Val Thr Gly Ala Ala
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Gly Gly Val Leu Leu Arg Pro Phe Ala Arg Leu Ile Ser Ser Thr Gly
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<213> Mycobacterium vaccae

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195

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200

Met Ala Lys Thr Ile Ala Tyr Asp Glu Glu Ala Arg Arg Gly Leu Glu

10

Gln Glu Ala Val Leu Glu Asp Pro Tyr Ile Leu Leu Val Ser Ser Lys 215 Val Ser Thr Val Lys Asp Leu Leu Pro Leu Leu Glu Lys Val Ile Gln 230 235 Ala Gly Lys Pro Leu Leu Ile Ile Ala Glu Asp Val Glu Gly Glu Ala 250 245 Leu Ser Thr Leu Val Val Asn Lys Ile Arg Gly Thr Phe Lys Ser Val 265 Ala Val Lys Ala Pro Gly Phe Gly Asp Arg Arg Lys Ala Met Leu Gln 280 Asp Met Ala Ile Leu Thr Gly Gly Gln Val Val Ser Glu Arg Val Gly 295 300 Leu Ser Leu Glu Thr Ala Asp Val Ser Leu Leu Gly Gln Ala Arg Lys 310 315 Val Val Val Thr Lys Asp Glu Thr Thr Ile Val Glu Gly Ser Gly Asp 325 330 Ser Asp Ala Ile Ala Gly Arg Val Ala Gln Ile Arg Ala Glu Ile Glu 345 Asn Ser Asp Ser Asp Tyr Asp Arg Glu Lys Leu Gln Glu Arg Leu Ala 360 Lys Leu Ala Gly Gly Val Ala Val Ile Lys Ala Gly Ala Ala Thr Glu 375 380 Val Glu Leu Lys Glu Arg Lys His Arg Ile Glu Asp Ala Val Arg Asn 390 395 Ala Lys Ala Ala Val Glu Glu Gly Ile Val Ala Gly Gly Val Ala 410 Leu Leu Gln Ser Ala Pro Ala Leu Asp Asp Leu Gly Leu Thr Gly Asp 425 Glu Ala Thr Gly Ala Asn Ile Val Arg Val Ala Leu Ser Ala Pro Leu 440 Lys Gln Ile Ala Phe Asn Gly Gly Leu Glu Pro Gly Val Val Ala Glu 455 460 Lys Val Ser Asn Leu Pro Ala Gly His Gly Leu Asn Ala Ala Thr Gly 470 475 Glu Tyr Glu Asp Leu Leu Lys Ala Gly Val Ala Asp Pro Val Lys Val 490 Thr Arg Ser Ala Leu Gln Asn Ala Ala Ser Ile Ala Ala Leu Phe Leu 505 500 Thr Thr Glu Ala Val Val Ala Asp Lys Pro Glu Lys Ala Ser Ala Pro 520 Ala Gly Asp Pro Thr Gly Gly Met Gly Gly Met Asp Phe 530 535

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Ile Ile Ala Glu Asp Val Glu Gly Glu Ala Leu Ser Thr Leu Val Val 35 40 45
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Asn Lys Ile Arg Gly Thr Phe Lys Ser Val Ala Val Lys Ala Pro Gly 50 55 60

Phe Gly Asp Arg Lys Ala Met Leu Gln Asp Met Ala Ile Leu Thr
65 70 75 80

Cly Cly Cly Cly Yel Yel Ser Cly Arg Yel Cly Ley Ser Ley Cly Thr Ala

Gly Gly Gln Val Val Ser Glu Arg Val Gly Leu Ser Leu Glu Thr Ala 85 90 95 Asp Val Ser Leu Leu Gly Gln Ala Arg Lys Val Val Val Thr Lys Asp

100 105 110

Glu Thr Thr Ile Val Glu Gly Ser Gly Asp Ser Asp Ala Ile Ala Gly

115 120 125 Arg Val Ala Gln Ile Arg Ala Glu Ile Glu Asn Ser Asp Ser Asp Tyr

130 135 140
Asp Arg Glu Lys Leu Gln Glu Arg Leu Ala Lys Leu Ala Gly Gly Val

145 150 155 160 Ala Val Ile Lys Ala Gly Ala Ala Thr Glu Val Glu Leu Lys Glu Arg

165 170 175
Lys His Arg Ile Glu Asp Ala Val Arg Asn Ala Lys Ala Ala Val Glu

180 185 190 Glu Gly Ile Val Ala Gly Gly Gly Val Ala Leu Leu Gln Ser Ala Pro

195 200 205 Ala Leu Asp Asp Leu Gly Leu Thr Gly Asp Glu Ala Thr Gly Ala Asn 210 215 220

Ile Val Arg Val Ala Leu Ser Ala Pro Leu Lys Gln Ile Ala Phe Asn 225 230 235 240

Gly Gly Leu Glu Pro Gly Val Val Ala Glu Lys Val Ser Asn Leu Pro 245 250 255

Ala Gly His Gly Leu Asn Ala Ala Thr Gly Glu Tyr Glu Asp Leu Leu 260 265 270

Lys Ala Gly Val Ala Asp Pro Val Lys Val Thr Arg Ser Ala Leu Gln 275 280 285

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660

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 Ala

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 Leu
 Arg
 Val
 Ser
 Asp
 Trp
 Pro
 Leu
 Tyr
 Met
 Ala
 Asp
 Gly

 Phe
 Ile
 Ala
 Ala
 Phe
 Gln
 Thr
 Ala
 Ser
 Gly
 Ile
 Thr
 Val
 Asp
 Tyr
 Lys

 Glu
 Asp
 Phe
 Asn
 Asp
 Asn
 Glu
 Gln
 Trp
 Phe
 Ala
 Lys
 Val
 Lys
 Glu
 Pro

 Glu
 Asp
 Asp
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 Gly
 Ala
 Asp
 Leu
 Val
 Ile
 Pro
 Thr
 Glu

 Fer
 Arg
 Lys
 Gln
 Asp
 Ile
 Ala
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 Leu
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 Ile
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 Thr
 Glu
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 Ile
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 Ile
 Asp
 Ile
 Asp
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145 150 155 160

Phe Ser Asp Val Gla Asp Gly Leu Gly Met Lle Met Leu Ser Gla Gly

Phe Ser Asp Val Gln Asp Gly Leu Gly Met Ile Met Leu Ser Gln Gly 165 170 175 Asn Ser Pro Glu Asn Pro Thr Thr Glu Ser Ile Gln Gln Ala Val Asp

180 185 190
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Leu Val Arg Glu Gln Asn Asp Arg Gly Gln Ile Arg Arg Phe Thr Gly

195

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205

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			180	-	-			185					Gly 190	_	
		195	_				200			_	_	205	Thr		
_	210	_	_			215					220		Leu		_
225					230					235			Arg		240
				245				-	250		_	_	Val	255	
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			340					345					350 Arg		
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			_	405	_			_	410				Val	415	
			420					425					430 Pro		
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<213> Mycobacterium vaccae

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60

Leu Ala Gly Ala Ser Phe Thr Asn Tyr Ser Arg Pro Val Gly Glu His

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Arg Leu Thr Val Val Thr Thr Phe Asn Ala Ala Asp Thr Pro Asp Asp
Val Cys Glu Met Leu Ser Ser Val Ala Ala Ser Leu Pro Glu Leu Arg
                85
                                    90
Thr Asp Gly Gln Ile Ala Thr Leu Tyr Leu Gly Ala Ala Glu Tyr Glu
                               105
Lys Ser Ile Pro Leu His Thr Pro Ala Val Asp Asp Ser Val Arg Ser
                            120
Thr Tyr Leu Arg Trp Val Trp Tyr Ala Ala Arg Arg Gln Glu Leu Arg
                       135
                                            140
Xaa Asn Gly Val Ala Asp Xaa Phe Asp Thr Pro Glu Arg Ile Ala Ser
                    150
                                        155
Ala Met Arg Ala Val Ala Ser Thr Leu Arg Leu Ala Asp Asp Glu Gln
                                    170
Gln Glu Ile Ala Asp Val Val Arg Leu Val Arg Tyr Gly Asn Gly Glu
                                185
Arg Leu Gln Gln Pro Gly Gln Val Pro Thr Gly Met Arg Phe Ile Val
                            200
Asp Gly Arg Val Ser Leu Ser Val Ile Asp Gln Asp Gly Asp Val Ile
                       215
Pro Ala Arg Val Leu Glu Arg Gly Asp Phe Leu Gly Gln Thr Thr Leu
                                       235
                   230
Thr Arg Glu Pro Val Leu Ala Thr Ala His Ala Leu Glu Glu Val Thr
                                    250
Val Leu Glu Met Ala Arg Asp Glu Ile Glu Arg Leu Val His Arg Lys
                                265
Pro Ile Leu Leu His Val Ile Gly Ala Val Ala Asp Arg Arg Ala His
                            280
Glu Leu Arg Leu Met Asp Ser Gln Asp
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     <213> Mycobacterium vaccae
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Leu Thr Asp Ile Arg Glu Ser Gln Ser Arg Gly Leu Glu Asn Gln Phe
                                25
Ala Asp Leu Lys Asn Ser Met Val Ile Tyr Ser Arg Gly Ser Thr Ala
                            40
Thr Glu Ala Ile Gly Ala Phe Ser Asp Gly Phe Arg Gln Leu Gly Asp
                        55
Ala Thr Ile Asn Thr Gly Gln Ala Ala Ser Leu Arg Arg Tyr Tyr Asp
                                        75
Arg Thr Phe Ala Asn Thr Thr Leu Asp Asp Ser Gly Asn Arg Val Asp
                85
                                    90
Val Arg Ala Leu Ile Pro Lys Ser Asn Pro Gln Arg Tyr Leu Gln Ala
                                105
Leu Tyr Thr Pro Pro Phe Gln Asn Trp Glu Lys Ala Ile Ala Phe Asp
                            120
Asp Ala Arg Asp Gly Ser Ala Trp Ser Ala Ala Asn Ala Arg Phe Asn
                        135
                                            140
Glu Phe Phe Arg Glu Ile Val His Arg Phe Asn Phe Glu Asp Leu Met
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Leu Leu Asp Leu Glu Gly Asn Val Val Tyr Ser Ala Tyr Lys Gly Pro Asp Leu Gly Thr Asn Ile Val Asn Gly Pro Tyr Arg Asn Arg Glu Leu Ser Glu Ala Tyr Glu Lys Ala Val Ala Ser Asn Ser Ile Asp Tyr Val Gly Val Thr Asp Phe Gly Trp Tyr Leu Pro Ala Glu Glu Pro Thr Ala Trp Phe Leu Ser Pro Val Gly Leu Lys Asp Arg Val Asp Gly Val Met Ala Val Gln Phe Pro Ile Ala Arg Ile Asn Glu Leu Met Thr Ala Arg Gly Gln Trp Arg Asp Thr Gly Met Gly Asp Thr Gly Glu Thr Ile Leu Val Gly Pro Asp Asn Leu Met Arg Ser Asp Ser Arg Leu Phe Arg Glu Asn Arq Glu Lys Phe Leu Ala Asp Val Val Glu Gly Gly Thr Pro Pro Glu Val Ala Asp Glu Ser Val Asp Arg Arg Gly Thr Thr Leu Val Gln Pro Val Thr Thr Arg Ser Val Glu Glu Ala Gln Arg Gly Asn Thr Gly Thr Thr Ile Glu Asp Asp Tyr Leu Gly His Glu Ala Leu Gln Ala Tyr Ser Pro Val Asp Leu Pro Gly Leu His Trp Val Ile Val Ala Lys Ile Asp Thr Asp Glu Ala Phe Ala Pro Val Ala Gln Phe Thr Arg Thr Leu Val Leu Ser Thr Val Ile Ile Ile Phe Gly Val Ser Leu Ala Ala Met Leu Leu Ala Arg Leu Phe Val Arg Pro Ile Arg Arg Leu Gln Ala Gly Ala Gln Gln Ile Ser Gly Gly Asp Tyr Arg Leu Ala Leu Pro Val Leu Ser Arg Asp Glu Phe Gly Asp Leu Thr Thr Ala Phe Asn Asp Met Ser Arg Asn Leu Ser Ile Lys Asp Glu Leu Leu Gly Glu Glu Arg Ala Glu Asn Gln Arg Leu Met Leu Ser Leu Met Pro Glu Pro Val Met Gln Arg Tyr Leu Asp Gly Glu Glu Thr Ile Ala Gln Asp His Lys Asn Val Thr Val Ile Phe Ala Asp Met Met Gly Leu Asp Glu Leu Ser Arg Met Leu Thr Ser Glu Glu Leu Met Val Val Val Asn Asp Leu Thr Arg Gln Phe Asp Ala Ala Ala Glu Ser Leu Gly Val Asp His Val Arg Thr Leu His Asp Gly Tyr Leu Ala Ser Cys Gly Leu Gly Val Pro Arg Leu Asp Asn Val Arg Arg Thr Val Asn Phe Ala Ile Glu Met Asp Arg Ile Ile Asp Arg His Ala Ala Glu Ser Gly His Asp Leu Arg Leu Arg Ala Gly Ile Asp Thr Gly Ser Ala Ala Ser Gly Leu Val Gly Arg Ser Thr Leu Ala Tyr Asp Met Trp Gly Ser Ala Val Asp Val Ala Asn Gln Val Gln Arg

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620
    610
                        615
Gly Ser Pro Gln Pro Gly Ile Tyr Val Thr Ser Arg Val His Glu Val
                    630
                                         635
Met Gln Glu Thr Leu Asp Phe Val Ala Ala Gly Glu Val Val Gly Glu
                                     650
Arg Gly Val Glu Thr Val Trp Arg Leu Gln Gly His Arg Arg
            660
                                665
                                                     670
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      <211> 520
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 179
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cccgacgacc tggcgtcgat tcgtaccggc cgcgcgaacc ccggcatgtt caaccggatc
                                                                       120
aacatcqact actacqqcqc ctccaccccq atcacqcaqc tqtccaqcat caacqtqccc
                                                                       180
gaggegegea tggtggtgat caagecetae gaggegagee agetgegeet categaggat
                                                                       240
gcgatccgca actccgacct cggcgtcaat ccgaccaacg acggcaacat catccgggtg
                                                                       300
tegatecege ageteacega ggagegeege egegacetgg teaageagge caaggecaag
                                                                       360
ggcgaggacg ccaaggtgtc ggtgcgcaac atccgtcgca acgatatgaa cacctttcgc
                                                                       420
ategeacegg taeggetgee gaegecaceg cegtegtaga agegacagag gategeaggt
                                                                       480
aacggtattg gccacgcctt ctgtggcggg ccgacaccac
                                                                       520
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      <211> 1071
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 180
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                                                                       120
teggeccage egggecteee geagecceeg etgecegeee etgecacagt gaegeaaace
                                                                       180
gtcacggttg cgcccaacgc cgcgccacaa ctcatcccgc gccccggtgt gacgcctgcc
                                                                       240
accggcggcg ccgccgcggt gcccgccggg gtgagcgccc cggcggtcgc gccggccccc
                                                                       300
gegetgeeeg eeegeeeggt gteeaegate geeeeggeea eetegggeae geteagegag
                                                                       360
ttettegeeg ceaagggegt cacgatggag cegeagteca geegegaett eegegeeete
                                                                       420
aacatcgtgc tgccgaagcc gcggggctgg gagcacatcc cggacccgaa cgtgccggac
                                                                       480
gcgttcgcgg tgctggccga ccgggtcggc ggcaacggcc tgtactcgtc gaacgcccag
                                                                       540
gtggtggtct acaaactcgt cggcgagttc gaccccaagg aagcgatcag ccacggcttc
                                                                       600
gtegacagee agaagetgee ggegtggegt tecacegaeg egtegetgge egacttegge
                                                                       660
ggaatgccgt cctcgctgat cgagggcacc taccgcgaga acaacatgaa gctgaacacg
                                                                       720
teceggegee aegteattge cacegegggg ceegaceact acetggtgte getgteggtg
                                                                       780
accaccageg tegaacagge egtggeegaa geegeggagg ceaeegaege gattgteaae
                                                                       840
ggetteaagg teagegttee gggteegggt eeggeegeac egecacetge acceggtgee
                                                                       900
cceggtgtcc cgcccgcccc cggcgccccg gcgctgccgc tggccgtcgc accacccccg
                                                                       960
getecegetg ttecegeegt ggegeeegeg ceacagetge tgggaetgea gggatagaeg
                                                                      1020
tcgtcgtccc ccgggcgaag cctggcgccc gggggacgac ggcccctttc t
                                                                      1071
      <210> 181
      <211> 152
      <212> PRT
      <213> Mycobacterium vaccae
      <400> 181
Val Ile Asp Glu Thr Leu Phe His Ala Glu Glu Lys Met Glu Lys Ala
                                    10
 1
                 5
                                                         15
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Val Ser Val Ala Pro Asp Asp Leu Ala Ser Ile Arg Thr Gly Arg Ala 25 Asn Pro Gly Met Phe Asn Arg Ile Asn Ile Asp Tyr Tyr Gly Ala Ser Thr Pro Ile Thr Gln Leu Ser Ser Ile Asn Val Pro Glu Ala Arg Met Val Val Ile Lys Pro Tyr Glu Ala Ser Gln Leu Arg Leu Ile Glu Asp 70 75 Ala Ile Arg Asn Ser Asp Leu Gly Val Asn Pro Thr Asn Asp Gly Asn 85 90 Ile Ile Arg Val Ser Ile Pro Gln Leu Thr Glu Glu Arg Arg Asp 105 100 Leu Val Lys Gln Ala Lys Ala Lys Gly Glu Asp Ala Lys Val Ser Val 120 Arg Asn Ile Arg Arg Asn Asp Met Asn Thr Phe Arg Ile Ala Pro Val 135 Arg Leu Pro Thr Pro Pro Pro Ser 145 150

<210> 182 <211> 331

<212> PRT

<213> Mycobacterium vaccae

<400> 182

Met Ser Glu Ile Ala Arg Pro Trp Arg Val Leu Ala Gly Gly Ile Gly 1 5 10 15

Ala Cys Ala Ala Gly Ile Ala Gly Val Leu Ser Ile Ala Val Thr Thr 20 25 30

Ala Ser Ala Gln Pro Gly Leu Pro Gln Pro Pro Leu Pro Ala Pro Ala

35 40 45

Thr Val Thr Gln Thr Val Thr Val Ala Pro Asn Ala Ala Pro Gln Leu 50 55 60 Ile Pro Arg Pro Gly Val Thr Pro Ala Thr Gly Gly Ala Ala Ala Val

65 70 75 80
Pro Ala Gly Val Ser Ala Pro Ala Val Ala Pro Ala Pro Ala Leu Pro

85 90 95
Ala Arg Pro Val Ser Thr Ile Ala Pro Ala Thr Ser Gly Thr Leu Ser

Glu Phe Phe Ala Ala Lys Gly Val Thr Met Glu Pro Gln Ser Ser Arg 115 120 125

Asp Phe Arg Ala Leu Asn Ile Val Leu Pro Lys Pro Arg Gly Trp Glu 130 135 140

Arg Val Gly Gly Asn Gly Leu Tyr Ser Ser Asn Ala Gln Val Val
165 170 175

Tyr Lys Leu Val Gly Glu Phe Asp Pro Lys Glu Ala Ile Ser His Gly
180 185 190

Phe Val Asp Ser Gln Lys Leu Pro Ala Trp Arg Ser Thr Asp Ala Ser 195 200 205

Leu Ala Asp Phe Gly Gly Met Pro Ser Ser Leu Ile Glu Gly Thr Tyr 210 215 220

Arg Glu Asn Asn Met Lys Leu Asn Thr Ser Arg Arg His Val Ile Ala 225 230 235 240

Thr Ala Gly Pro Asp His Tyr Leu Val Ser Leu Ser Val Thr Thr Ser 245 250 255

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Val Glu Gln Ala Val Ala Glu Ala Glu Ala Thr Asp Ala Ile Val
Asn Gly Phe Lys Val Ser Val Pro Gly Pro Gly Pro Ala Ala Pro Pro
                            280
Pro Ala Pro Gly Ala Pro Gly Val Pro Pro Ala Pro Gly Ala Pro Ala
                                            300
                        295
Leu Pro Leu Ala Val Ala Pro Pro Pro Ala Pro Ala Val Pro Ala Val
                                        315
Ala Pro Ala Pro Gln Leu Leu Gly Leu Gln Gly
                325
      <210> 183
      <211> 207
      <212> DNA
      <213> Mycobacterium vaccae
      <400> 183
acctacgagt tcgagaacaa ggtcacgggc ggccgcatcc cgcgcgagta catcccgtcg
                                                                        60
gtggatgccg gcgcgcagga cgccatgcag tacggcgtgc tggccggcta cccgctggtt
                                                                       120
aacgtcaagc tgacgctgct cgacggtgcc taccacgaag tcgactcgtc ggaaatggca
                                                                       180
ttcaaggttg ccggctccca ggtcata
                                                                       207
      <210> 184
      <211> 69
      <212> PRT
      <213> Mycobacterium vaccae
      <400> 184
Thr Tyr Glu Phe Glu Asn Lys Val Thr Gly Gly Arg Ile Pro Arg Glu
                 5
Tyr Ile Pro Ser Val Asp Ala Gly Ala Gln Asp Ala Met Gln Tyr Gly
                                25
Val Leu Ala Gly Tyr Pro Leu Val Asn Val Lys Leu Thr Leu Leu Asp
                            40
Gly Ala Tyr His Glu Val Asp Ser Ser Glu Met Ala Phe Lys Val Ala
                        55
                                            60
Gly Ser Gln Val Ile
      <210> 185
      <211> 898
      <212> DNA
      <213> Mycobacterium vaccae
      <220>
      <221> unsure
      <222> (637)...(637)
      <221> unsure
      <222> (662)...(662)
      <400> 185
cgacctccac ccgggcgtga ggccaaccac taggctggtc accagtagtc gacggcacac
                                                                        60
ttcaccgaaa aaatgaggac agaggagaca cccgtgacga tccgtgttgg tgtgaacggc
                                                                       120
tteggeegta teggaegeaa ettetteege gegetggaeg egeagaagge egaaggeaag
                                                                       180
aacaaggaca tegagategt egeggteaac gaceteaceg acaaegeeac getggegeac
                                                                       240
ctgctgaagt tcgactcgat cctgggccgg ctgccctacg acgtgagcct cgaaggcgag
                                                                       300
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gacaccatcg tcgtcggcag caccaagatc aaggcgctcg aggtcaagga aggcccggcg
gegetgeect ggggegacet gggegtegae gtegtegteg agtecacegg catetteace
aagegegaca aggeecaggg ecacetegac gegggegeca agaaggteat cateteegeg
ceggecaceg atgaggacat caccategtg eteggegtea aegaegacaa gtaegaegge
agccagaaca tcatctccaa cgcgtcgtgc accacgaact gcctcggccc gctggcgaag
gtcatcaacg acgagttcgg catcgtcaag ggcctgntga ccaccatcca cgcctacacc
enggtecaga acetgeagga eggeeegeac aaggatetge geegggeeeg egeegeege
ctgaacatcg tgccgacctc caccggtgcc gccaaggcca tcggactggt gctgcccgag
ctgaagggca agctcgacgg ctacgcgctg cgggtgccga tccccaccgg ctcggtcacc
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      <210> 186
      <211> 268
      <212> PRT
      <213> Mycobacterium vaccae
      <220>
      <221> UNSURE
      <222> (182)...(182)
      <221> UNSURE
      <222> (190)...(190)
      <400> 186
Val Thr Ile Arg Val Gly Val Asn Gly Phe Gly Arg Ile Gly Arg Asn
                 5
Phe Phe Arq Ala Leu Asp Ala Gln Lys Ala Glu Gly Lys Asn Lys Asp
            20
Ile Glu Ile Val Ala Val Asn Asp Leu Thr Asp Asn Ala Thr Leu Ala
                            40
His Leu Leu Lys Phe Asp Ser Ile Leu Gly Arg Leu Pro Tyr Asp Val
                        55
Ser Leu Glu Gly Glu Asp Thr Ile Val Val Gly Ser Thr Lys Ile Lys
                    70
                                         75
Ala Leu Glu Val Lys Glu Gly Pro Ala Ala Leu Pro Trp Gly Asp Leu
                85
                                    90
Gly Val Asp Val Val Val Glu Ser Thr Gly Ile Phe Thr Lys Arg Asp
            100
                                105
                                                     110
Lys Ala Gln Gly His Leu Asp Ala Gly Ala Lys Lys Val Ile Ile Ser
                            120
Ala Pro Ala Thr Asp Glu Asp Ile Thr Ile Val Leu Gly Val Asn Asp
                        135
                                            140
Asp Lys Tyr Asp Gly Ser Gln Asn Ile Ile Ser Asn Ala Ser Cys Thr
                    150
                                         155
Thr Asn Cys Leu Gly Pro Leu Ala Lys Val Ile Asn Asp Glu Phe Gly
                165
                                     170
Ile Val Lys Gly Leu Xaa Thr Thr Ile His Ala Tyr Thr Xaa Val Gln
            180
                                185
                                                     190
Asn Leu Gln Asp Gly Pro His Lys Asp Leu Arg Arg Ala Arg Ala Ala
                            200
Ala Leu Asn Ile Val Pro Thr Ser Thr Gly Ala Ala Lys Ala Ile Gly
    210
                        215
Leu Val Leu Pro Glu Leu Lys Gly Lys Leu Asp Gly Tyr Ala Leu Arg
                    230
                                         235
Val Pro Ile Pro Thr Gly Ser Val Thr Asp Leu Thr Ala Glu Leu Gly
                245
                                     250
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420

480

540

600

660

720

780

840

898

Lys Ser Ala Thr Val Asp Glu Ile Asn Ala Ala Met

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260
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      <211> 41
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      <213> Mycobacterium vaccae
      <220>
      <221> UNSURE
      <222> (39)...(39)
      <400> 187
Met Asn Lys Ala Glu Leu Ile Asp Val Leu Thr Glu Lys Leu Gly Ser
Asp Arg Arg Gln Ala Thr Ala Ala Val Glu Asn Val Val Asp Thr Ile
            20
                                 25
Val Ala Ala Val Pro Lys Xaa Val Val
      <210> 188
      <211> 26
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <221> unsure
      <222> (12)...(12)
      <400> 188
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                                                                         26
      <210> 189
      <211> 20
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <400> 189
                                                                         20
atsgtrtgva cvacgttytc
      <210> 190
      <211> 84
      <212> DNA
      <213> Artificial Sequence
      <220>
      <223> Made in a lab
      <221> unsure
      <222> (2)...(2)
      <400> 190
                                                                         60
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atggagaacg tggtccacac cata	84
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<220> <221> unsure <222> (2)(2)	
<400> 191	
gnactcattg acgtactcac tgagaagctg ggctcggatt gtcggcaagc gactgcggcg gtggagaatg ttgtcgacac catcgtgcgc gccgtgcaca agggtgagag cgtcaccatc acgggcttcg gtgttttcga gcagcgtcgt cgcgcagcac gcgtggcacg caatccgcgc accggcgaga ccgtgaaggt caagcccacc tcagtcccgg cattccgtcc cggcgctcag ttcaaggctg ttgtctctgg cgcacagaag cttccggccg agggtccggc ggtcaagcgc ggtgtgaccg cgacgagcac cgcccgcaag gcagcca	60 120 180 240 300 337
<210> 192 <211> 111 <212> PRT <213> Mycobacterium vaccae	
<220> <221> UNSURE <222> (1)(1)	
<400> 192	
Xaa Leu Ile Asp Val Leu Thr Glu Lys Leu Gly Ser Asp Arg Gln Ala 1 5 10 15	
Thr Ala Ala Val Glu Asn Val Val Asp Thr Ile Val Arg Ala Val His 20 25 30	
Lys Gly Glu Ser Val Thr Ile Thr Gly Phe Gly Val Phe Glu Gln Arg 35 40 45	
Arg Arg Ala Ala Arg Val Ala Arg Asn Pro Arg Thr Gly Glu Thr Val 50 55 60	
Lys Val Lys Pro Thr Ser Val Pro Ala Phe Arg Pro Gly Ala Gln Phe 65 70 75 80	
Lys Ala Val Val Ser Gly Ala Gln Lys Leu Pro Ala Glu Gly Pro Ala 85 90 95	
Val Lys Arg Gly Val Thr Ala Thr Ser Thr Ala Arg Lys Ala Ala 100 105 110	
<210> 193 <211> 1164 <212> DNA <213> Mycobacterium vaccae	
<400> 193	
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geggeggeea getggtaegg ceattecage gtgetgateg aggtegaegg etacegegtg	420

ctggccgacc	cggtgtggag	caacagatgt	tcgccctcac	gggcggtcgg	accgcagcgc	480
atgcacgacg	tcccggtgcc	gctggaggcg	cttcccgccg	tggacgcggt	ggtgatcagc	540
cacgaccact	acgaccacct	cgacatcgac	accatcgtcg	cgttggcgca	cacccagcgg	600
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gcgcggatcg	tcgagttgga	ctggcacgaa	gcccaccgca	tagacgacct	gacgctggtc	720
tgcacccccg	cccggcactt	ctccggacgg	ttgttctccc	gcgactcgac	gctgtgggcg	780
tcgtgggtgg	tcaccggctc	gtcgcacaag	gcgttcttcg	gtggcgacac	cggatacacg	840
aagagcttcg	ccgagatcgg	cgacgagtac	ggtccgttcg	atctgaccct	gctgccgatc	900
ggggcctacc	atcccgcgtt	cgccgacatc	cacatgaacc	ccgaggaggc	ggtgcgcgcc	960
catctggacc	tgaccgaggt	ggacaacagc	ctgatggtgc	ccatccactg	ggcgacattc	1020
cgcctcgccc	cgcatccgtg	gtccgagccc	gccgaacgcc	tgctgaccgc	tgccgacgcc	1080
gagcgggtac	gcctgaccgt	gccgattccc	ggtcagcggg	tggacccgga	gtcgacgttc	1140
gacccgtggt	ggcggttctg	aacc				1164

<210> 194

<211> 370

<212> PRT

<213> Mycobacterium vaccae

5

<400> 194

1				2					TO					13	
Ala	Gly	Gly	Phe 20	Val	Leu	Arg	Ala	Leu 25	Gln	Gly	Thr	Pro	Ala 30	Ala	Leu
Gly	Ala	Thr 35	Pro	Gly	Glu	Val	Ala 40	Pro	Val	Ala	Arg	Arg 45	Ser	Pro	Asn
Tyr	Arg 50	Asp	Gly	Lys	Phe	Val 55	Asn	Leu	Glu	Pro	Pro 60	Ser	Gly	Ile	Thr
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Ala	Ser	Gln	Gly	Lys 85	Pro	Pro	Gly	Pro	Ile 90	Pro	Leu	Ala	Glu	Pro 95	Pro
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